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(71) Applicants: BAYLOR COLLEGE OF MEDICINE [US/US]; One Baylor Plaza, Houston, TX 77030 (US). UNIT-ED SATES OF AMERICA [US/US]; Department of Health and Human Services, National Institute of Health, Office of Technology Transfer, Bethesda, MD 20892 (US).

(72) Inventors: ROOP, Dennis, R.; 2331 Bellefontaine, Houston, TX 77030 (US). ROTHNAGEL, Joseph, A.; 5427 Queensloch Drive, Houston, TX 77096 (US). GREENHALGH, David, A.; 7423 Brompton, Houston, TX 77025 (US). YUSPA, Stuart, H.; 8013 Carita Court, Bethesda, MD 20817 (US).

(74) Agent: PAUL, Thomas, D.; Fulbright & Jaworski, 1301 McKinney, Suite 5100, Houston, TX 77010-3095 (US).

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(54) Title: DEVELOPMENT OF A VECTOR TO TARGET GENE EXPRESSION TO THE EPIDERMIS OF TRANSGENIC ANIMALS

(57) Abstract

The keratin K1 vector for expression of a nucleic acid cassette in the epidermis. The vector has a 5' flanking region of the keratin K1 gene including a 5' flanking sequence, a keratin K1 promoter, 5' transcribed but untranslated region and a first intron and an intron/exon boundary, all in sequential and positional relationship for expression of a nucleic acid cassette. A 3' flanking region in the K1 keratin gene containing regulatory sequences including a 3' transcribed but untranslated region and contiguous noncoding DNA containing the transcriptional termination region. The 3' flanking region and a 5' flanking region are linked by sette encodes a sequence which can be used to express any protein, polypeptide or antisense RNA. The vector can be inserted both in vivo and ex vivo into epidermal cells. Further, it can be used for making transgenic animals or bioreactors of epidermal cells. The vector has been found useful for gene therapy in treatment of a variety of diseases including skin ulcers, wound healing, surgical incisions, psoriasis and cancer.

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DEVELOPMENT OF A VECTOR TO TARGET GENE EXPRESSION TO THE EPIDERMIS OF TRANSGENIC ANIMALS

This invention was partially supported by grants from the United States government under HD25479, AI30283 and CA52607 awarded by the National Institutes of Health. Further, this work was partially performed at the National Institutes of Health in the Laboratory of Cellular Carcinogenesis and Tumor Promotion, Division of Cancer Etiology, National Cancer Institute. The government has certain rights in the invention.

FIELD OF THE INVENTION

The present invention relates to expression vectors for use in expressing polypeptides in epidermal cells of transgenic animals. More particularly it relates to vectors containing the K1 keratin gene promoter, its 5' flanking region, its 5' transcribed but untranslated region, its first intron and intron/exon boundary, its 3' transcribed but untranslated region, its contiguous non-coding DNA containing the gene's natural transcriptional termination region and its 3' flanking region.

BACKGROUND OF THE INVENTION

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The ability to stably introduce genes into the germline of mice has greatly enhanced prospects for the generation of animal models of human diseases (Palmiter and Brinster, Ann. Rev. Genet., Vol. 20, pp. 465-499 (1986)). The need for such animal models is becoming increasingly apparent as novel pharmaceuticals are developed which are specifically

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designed to inhibit expression of human viruses or counteract the effect of mutated genes that occur in human diseases. Current efficacy assessments of these new therapeutic agents are restricted to in vitro models which do not allow evaluation of delivery routes nor assessment of other factors known to affect disease processes in vivo, such as blood supply, an intact immune system, humoral and cell-mediated growth controls and physical barriers to disease progression. In addition, the prospects for utilizing gene therapy to treat human disorders are coming closer to reality. Therefore, animal models of human diseases would be useful to assess the therapeutic potential of these approaches. epidermis is an attractive tissue for the development animal models since it serves as a general model for other squamous epithelia and its accessibility allows macroscopic observation of pathological events and easy assessment of therapeutic potential. The development of a vector which specifically targets gene expression to the epidermis of transgenic animals is the subject of this invention.

The epidermis is a continuously regenerating stratified squamous epithelium. Differentiated epidermal cells are the progeny of proliferative cells located in the basal cell layer and there is substantial evidence suggesting that the regeneration process occurs in proliferative units composed of slowly cycling, self-renewing stem cells, proliferative but non-renewing transit amplifying cells, and post-mitotic maturing epidermal cells (Iversen, et al., Cell Tissue Kinet., Vol. 1, pp. 351-367, (1968); MacKenzie, et al., Nature, Vol. 226, pp. 653-655, (1970); Christophers, et al., J. Invest. Dermatol., Vol. 56, pp. 165-170, (1971); Potten, In Stem Cells: Their Identification and Characterization, pp. 200-232, (1983); Cotsarelis, et al., Cell, Vol. 61, pp. 1329-1337, (1990)). The maturation process (terminal differentiation) is initiated when epidermal cells withdraw from the cell cycle and migrate from the basal layer into the spinous layer. Maturation continues as spinous cells migrate into the

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granular layer and terminates with the formation of the stratum corneum. Morphological and biochemical studies have shown that terminal differentiation occurs in stages. (Matoltsy, J. Invest. Dermatol., Vol. 65, pp. 127-142, (1975)). Keratins K5 and K14 are major products of basal epidermal cells (Woodcock-Mitchell, et al., J. Cell Biol., Vol. 95, pp. 580-588, (1982)). These proteins assemble into 10 nm filaments (intermediate filaments [IF]) and, together with microtubules (tubulin) microfilaments (actin), comprise the cytoskeleton of epidermal cells (Steinert, P.M., et al., Cell, Vol. 42, pp. 411-419, (1985)). One of the earliest changes associated with the commitment to differentiation and migration into the spinous layer is the induction of another differentiation-specific pair of keratins (K1 and K10). IF containing K1 and K10 replace those containing K5 and K14 as the major products of cells in the spinous layer (Woodcock-Mitchell, et al., J. Cell Biol., Vol. 95, pp. 580-588, (1982); Roop, et al., Proc. Natl. Acad. Sci., USA, Vol. 80, pp. 716-720, (1983); Schweizer, et al, Cell, Vol. 37, pp. 159-170, (1984)). The keratin IF formed by these proteins assemble into bundles. In the granular layer, another high molecular weight non-IF protein is synthesized, which is processed into filaggrin, and is thought to promote keratin filament aggregation and disulfide-bond formation (Dale, B.A., et al., Nature, Vol. 276, pp. 729-731, (1978); Harding, C.R., et al., J. Mol. Biol., Vol. 170, pp. 651-673, (1983)). In the final stage of epidermal cell maturation, transglutaminase catalyzes the crosslinking of involucrin and loricrin, by the formation of $(\gamma$ -glutamyl) lysine isopeptides, into a highly insoluble cornified envelope which is located just beneath the plasma membrane (Rice and Green, Cell Vol. II, pp. 417-422 (1977) Mehrel, et al., Cell, Vol. 61, pp. 1103-1112, (1990)).

Genes or cDNAs encoding the major keratins expressed in epidermal cells have now been cloned: K5 (Lersch, et al., Mol. and Cell Biol., Vol. 8, pp. 486-493, (1988), K14 (Marchuk, et al., Proc. Natl. Acad.

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Sci, USA, Vol. 82, pp. 1609-1613, (1985); Knapp, et al., J. Biol. Chem, Vol. 262, pp. 938-945, (1987); Roop, et al., Cancer Res., Vol. 48, pp. 3245-3252, (1988), K1 (Steinert, et al., J. Biol. Chem., Vol. 260, pp. 7142-7149, (1985) and K10 (Krieg, et al., J. Biol. Chem., Vol. 260, pp. 5867-5870, (1985)). Northern blot analysis and in situ hybridization studies suggest that keratin genes K5 and K14 are predominantly transcribed in the proliferating basal layer and transcription of keratin genes K1 and K10 is induced as cells migrate into the spinous layer (Lersch, et al., Mol. and Cell Biol., Vol. 8, pp. 486-493, (1988); Knapp, et al., J. Biol. Chem., Vol. 262, pp. 938-945, (1987); Roop, et al., Cancer Res., Vol. 48, pp. 3245-3252, (1988)). Genes encoding rat (Haydock, et al., J. Biol. Chem., Vol. 261, pp. 12520-12525, (1986)) and mouse (Rothnagel, et al., J. Biol. Chem., Vol. 262, pp. 15643-15648, (1987)) filaggrin have now been identified and in situ hybridization experiments have confirmed that transcription of this gene is restricted to the granular layer (Rothnagel, et al, J. Biol. Chem., Vol. 262, pp. 15643-15648, (1987); Fisher, et al. J. Invest. Dermatol., Vol. 88, pp. 661-664, (1987)). To date, loricrin is the only gene encoding a component of the cornified envelope to be studied at the molecular level by in situ hybridization and transcripts of this gene are restricted to the granular layer (Mehrel, et al., Cell, Vol. 61, pp. 1103-1112, (1990)).

From this description of gene expression in the epidermis, there would appear to be many candidate genes from which to choose for targeting to the epidermis. However, this is not the case. Keratins K5 and K14, expressed in the proliferative compartment of the epidermis, are not only expressed in the epidermis but in all squamous epithelia. Furthermore, these genes are expressed early in development (Dale and Holbrook, In: Current Topics in Developmental Biology, pp. 127-151, (1987)) and this could cause lethality in utero. The generation of animal models of hyperproliferative diseases such as cancer and psoriasis would most likely requir expression in the basal compartment in cells with

proliferative potential, therefore, genes expressed post-mitotically such as keratins K1 and K10 and those encoding filaggrin and components of the cell envelope (involucrin and loricrin) would be excluded on this basis. The present invention, however, demonstrates that a 12 kb fragment of the human keratin gene K1 (HK1) contains sequences regulating tissue and developmental specific expression in transgenic mice. This fragment lacks sequences responsive to negative control of differentiation specific expression resulting in expression of the HK1 gene in some cells of the basal cell compartment of the epidermis. Although regulatory elements of the HK1 gene fail to completely mimic the expression pattern of the endogenous mouse K1 gene, they are ideally suited for targeting gene expression for the following reasons: (1) expression only occurs in the epidermis and not other squamous epithelia; (2) expression occurs at a late stage of development (day 15) and, therefore, is unlikely to result in lethality it utero; (3) expression occurs in a large proportion of basal cells that have proliferative potential.

SUMMARY OF THE INVENTION

An object of the present invention is a keratin K1 vector for expressing nucleic acid sequences in the epidermis.

An additional object of the present invention is a keratin K1 vector containing an oncogene.

A further object of the present invention is a bioreactor for producing proteins, polypeptides and antisense RNA in transduced epidermal cells.

An additional object of the present invention is an *in vivo* method of transducing epidermal cells with a keratin K1 vector.

A further object of the present invention is provision of a transgenic animal containing the keratin K1 epidermal vector.

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An additional object of the present invention is the provision of a transgenic animal for the study of cancer.

Another object of the present invention is a method of treating skin ulcers.

A further object of the present invention is an enhanced method of wound healing or healing of surgical incisions.

An additional object of the present invention is a method of treating psoriasis.

An additional object of the present invention is a method of treating skin cancer.

An additional object of the present invention is a vaccination procedure using the keratin K1 epidermal vector.

Thus, in accomplishing the foregoing objects, there is provided in accordance with one aspect of the present invention a keratin K1 vector for expression of a nucleic acid cassette in the epidermis comprising: a 5' flanking region of the keratin K1 gene, said 5' flanking sequence including a keratin K1 promoter, a 5' transcribed but untranslated region and a first intron and an intron/exon boundary all in sequential and positional relationship for expression of a nucleic acid cassette; a 3' flanking region of the keratin K1 gene containing regulatory sequences, said 3' flanking region including a 3' transcribed but untranslated region and contiguous noncoding DNA containing a transcriptional termination region; and a polylinker having a plurality of restriction endonuclease sites, said polylinker connecting the 5' flanking region to the 3' flanking region and said polylinker further providing a position for insertion of the nucleic acid cassette.

In specific embodiments of the present invention the keratin K1 vector has a 5' flanking region of approximately 1.2 kb and a 3' flanking region of approximately 3.9 kb.

In alternative embodiments of the present invention there is a further addition of approximately 8.0 kb of 5' flanking sequence from the 18 kb Eco RV fragment onto the end of the vector.

In another alternative embodiment the Vitamin D, regulatory element within the human K1 keratin gene is identified and utilized to suppress expression of the keratin K1 vector.

In the specific embodiments of the present invention the keratin K1 vector is used to transduce epidermal cells to form bioreactors. The bioreactors produce a variety of proteins, polypeptides and RNAs. Additionally, the vector can be used to form transgenic animals. The transgenic animals can be used to study cancer, drug reactions and treatments.

The keratin K1 vector can also be used for the treatment of a variety of diseases, including wounds, surgical incisions, psoriasis, skin ulcers and skin cancer and can be used for the production of vaccines.

Other and further objects, features and advantages will be apparent from the following description of the presently preferred embodiments of the invention which are given for the purposes of disclosure when taken in conjunction with the accompanying drawings.

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BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 is a schematic drawing of the human keratin K1 gene (HK1) and the expression vector derived from its regulatory sequences.

Figure 2 shows the expression characteristics of the HK1 vector in vivo in transgenic mice utilizing a reporter gene encoding E. coli β-galactosidase. Figure 3 demonstrates the suppression of the SV40 promoter by a novel negative regulatory element from the HK1 gene (HK1.NRE) in the

presence of Vitamin D_3 .

Figure 4 is a schematic drawing of the HK1 vector containing the coding sequence of v-ras^{Ha} protein of Harvey Murine Sarcoma Virus.

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Figure 5 is a schematic drawing of the HK1 vector containing the coding sequence of the v-fos protein from a FBJ/FBR chimeric plasmid.

Figure 6 is a schematic drawing of the HK1 expression vector containing the coding sequences of the E6 and E7 proteins from human papilloma virus 18.

Figure 7 is a schematic drawing of the HK1 vector containing the coding sequence of $TGF-\alpha$.

Figure 8 is a schematic drawing of the HK1 vector containing the coding sequence of the trans-regulatory protein *tat*, from human immunodeficiency virus.

Figure 9 is a schematic drawing of an 18kb Eco RV fragment containing the HK1 gene.

Figure 10 is a schematic drawing of a derivative of the HK1 vector containing additional 5' flanking sequences which restrict expression to differentiated epidermal cells.

The drawings are not necessarily to scale, and certain features of the invention may be exaggerated in scale and shown in schematic form in the interest of clarity and conciseness.

DETAILED DESCRIPTION OF THE INVENTION

It will be readily apparent to one skilled in the art that varying substitutions and modifications may be made to the invention disclosed herein without departing from the scope and spirit of the invention.

The term "transformed" as used herein refers to the process or mechanism of inducing changes in the characteristics (expressed phenotype) of a cell by the mechanism of gene transfer whereby DNA is introduced into a cell in a form where it expresses a specific gene product or alters expression of endogenous gene products.

The term "transduction" as used herein refers to the process of introducing a DNA expression vector into a cell. Various methods of

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transduction are possible, including microinjection, CaPO₄, lipofection (lysosome fusion), use of a gene gun and DNA vector transporter. The human keratin K1 vector can be transduced into the epidermal cells by any of the variety of ways described above.

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The term "DNA vector transporter" as used herein refers to those molecules which bind to DNA vectors and are capable of being taken up by epidermal cells. DNA transporter is a molecular complex capable of non-covalent binding to DNA and efficiently transporting the DNA through the cell membrane. Although not necessary, it is preferable that the transporter also transport the DNA through the nuclear membrane.

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The term "transient" as used in transient transfection, transient transduction or transiently transformed relates to the introduction of genes into the epidermal cells to express specific proteins, polypeptides and RNA wherein the introduced genes are not integrated into the host cell genome and accordingly are eliminated from the cell over a period of time. Transient expression relates to the expression of gene products during the period of transient transfection. Additionally, transient can refer to a stable transfection or transduction into cells, where the cells die and are sloughed off from the skin. Thus, the transformed cells are only transiently available for the expression of the incorporated genes.

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The term "stable" as used in stable transfection, stable transduction or stably transformed refers to the introduction of genes into the chromosome of the targeted cell where it integrates and becomes a permanent component of the genetic material in that cell. Gene expression after stable transduction can permanently alter the characteristics of the cell leading to stable transformation. An episomal transformation is a variant of stable transformation in which the introduced gene is not incorporated into the host cell chromosomes but rather is replicated as an extrachromosomal element. This can lead to apparently stable transformation of the characteristics of the cell. As

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indicated above, in epidermal cells, which are sloughed from the body through the skin, stable transformation can become transient transformation because the cells are lost.

The term "nucleic acid cassette" as used herein refers to the genetic material of interest which can express a protein, polypeptide or RNA and which is capable of being incorporated into the epidermal cells. The nucleic acid cassette is positionally and sequentially oriented within the keratin K1 vector such that the nucleic acid in the cassette can be transcribed into RNA or antisense RNA and, when necessary, translated into proteins or polypeptides in the transformed epidermal cells. A variety of proteins and polypeptides can be expressed in the transformed epidermal cells by the sequence in the nucleic acid cassette. proteins or polypeptides which can be expressed include hormones, growth factors, enzymes, clotting factors, apolipoproteins, receptors, drugs, tumor antigens, viral antigens, parasitic antigens, bacterial antigens and oncogenes. Specific examples of these compounds include proinsulin, insulin, growth hormone, insulin-like growth factor I, insulin-like growth factor II, insulin growth factor binding protein, epidermal growth factor TGF-α, dermal growth factor PDGF, angiogenesis factor, for instance, acid fibroblast and basic fibroblast growth factors and angiogenin, matrix protein, such as, Type IV collagen, Type VII collagen, laminin, nidogen and proteins from viral, bacterial and parasitic organisms which can be used to induce an immunologic response.

In addition, the nucleic acid cassette can encode a "transforming gene" which encompasses viral oncogenes, endogenous proto-oncogenes and activated proto-oncogenes. A variety of oncogenes are known in the art. The term "oncogene" means those genes which cause cancer and include both viral and cellular oncogenes, many of which are homologous to DNA sequences endogenous to rodents and/or humans. The term

oncogene includes both the viral sequence and the homologous endogenous sequences. Some examples of transforming genes are listed in Table 1.

Table 1.
Transforming Genes

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ABBREVIATION	NAME
Ha-ras	Harvey Murine Sarcoma Virus
Ki-ras	Kirsten Murine Sarcoma Virus
N-ras	Neuroblastoma oncogene
fos	FBJ or FBR osteosarcoma virus
myc	Avian MC29 myelocytomatosis virus
src	Rous sarcoma virus
sis	Simian sarcoma virus/PDGF β chain
erbA	Avian erythroblastosis virus/Thyroxine T3 receptor
erbB	Avian erythroblastosis virus/Truncated EGF receptor
jun	Avian sarcoma virus 17
p Large T	Polyomavirus transforming gene
p Middle T	Polyomavirus transforming gene
HPV E7	Early region transforming gene from human papillomavirus 6, 11, 16, 18
HPV E6	Early region transforming gene from human papilloma virus 6, 11, 16, 18

HPV E5	Early region transforming gene from human papilloma virus 6, 11, 16, 18
tat	HIV transforming gene
EIA	Adenovirus early region 1A
Rb	Mutated retinoblastoma gene
p53	Mutated p53 anti-oncogene
WT1	Mutated Wilms tumor anti-oncogene
TGF-α	Transforming growth factor α
TGF-β	Transforming growth factor β
EGFR	Mutated epidermal growth factor receptor
RAR	Mutated retinoic acid receptor
VD _s R	Mutated vitamin D ₃ receptor
PKC	Mutated protein kinase C

The genetic material which is incorporated into the epidermal cells using the keratin K1 vector includes DNA not normally found in epidermal cells, DNA which is normally found in epidermal cells but not expressed at physiologically significant levels, DNA normally found in epidermal cells and normally expressed at physiological desired levels, any other DNA which can be modified for expression in epidermal cells, and any combination of the above.

The term "keratin K1 vector" or "HK1 vector" as used herein is a vector which is useful for expression of a nucleic acid sequence in epidermal cells. The keratin K1 vector comprises a 5' flanking region of

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the keratin K1 gene, said flanking region including a promoter, a first intron and an intron/exon boundary all in sequential and positional relationship for the expression of a nucleic acid cassette; a 3' flanking sequence of a keratin K1 gene; and a poly-linker. The poly-linker includes a plurality of restriction endonuclease sites. The polylinker connects the 5' flanking region to the 3' flanking sequence and further provides a position for insertion of the nucleic acid cassette.

The sequence for the 3' flanking region of the human keratin K1 gene contains regulator elements and is used for preparing the keratin K1 vector. It is shown in SEQ. ID No. 1. The keratin K1 vector has a 5' flanking region comprising nucleotides 1 to 1246 of SEQ. ID. No. 1; a 3' flanking sequence containing regulatory sequences comprises nucleotides 6891 to 10747 of SEQ. ID. No. 1; and a poly linker comprising nucleotides 2351 to 2376 of SEQ. ID. No. 2 (the HK1 expression vector).

The keratin K1 vector has a 5' flanking region of approximately 1.2 kb, an intron and intron/exon boundary of approximately 1.0 kb and a 3' flanking sequence of approximately 3.9 kb.

The restriction endonuclease sites found in the linker and polylinker of the keratin K1 vector can be any restriction endonucleases which will allow insertion of the nucleic acid cassette. In the preferred embodiment they are usually selected from the group consisting of Bam HI, Kpn I, Cla I, Not I, Xma I, and Bgl II.

One skilled in the art will readily recognize that there are a variety of ways to introduce the keratin K1 vector into epidermal cells. The vectors can be inserted either in vivo or ex vivo. The mode of insertion will, to a certain degree, determine the available methods for the insertion. The in vivo insertion is preferred for gene therapy. In this procedure the human keratin K1 vector is contacted with epidermal cells for sufficient time to transform the epidermal cells.

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One embodiment of the present invention includes a bioreactor. A bioreactor is comprised of transformed epidermal cells which contain the keratin K1 vector. Once the vector is inserted in the epidermal cells, the epidermal cells will express the nucleic cassette and produce the protein, polypeptide or antisense RNA of interest. This can be done either *in vivo* or *ex vivo*. Any compound which can be encoded in, and expressed by, the nucleic acid cassette can be produced by the bioreactor.

One method for ex vivo introduction of the keratin K1 vector into epidermal cells includes a cotransfection of the vector with a selectable marker. The selectable marker is used to select those cells which have become transformed. The cells can then be used in any of the methods described in the present invention.

Another embodiment of the present invention is a method of making transgenic animals comprising the steps of inserting the human keratin K1 vector into the embryo of the animal. The transgenic animal can include the resulting animal in which the vector has been inserted into the embryo or any progeny. The term progeny as used herein includes direct progeny of the transgenic animal as well as any progeny of succeeding progeny. Thus, one skilled in the art will readily recognize that if two different transgenic animals have been made using different genes in the nucleic acid cassette and they are mated, the possibility exists that some of the resulting progeny will contain two or more introduced sequences. One skilled in the art will readily recognize that by controlling the matings, transgenic animals with multiple vectors can be made.

In the transgenic animals that contain the human keratin K1 vector in its germ and somatic cells, the nucleic acid cassette of the said vector is only expressed in the epidermal cells. This is a distinct advantage over other transgenic animal models where there is not as much control over the expression of the sequence in the tissues.

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In the preferred embodiment, the transgenic animal will contain an oncogene sequence in the nucleic acid cassette. Preferably the animal is a rodent. The transgenic animal can be used in any method for studying a variety of diseases including the origin of cancer, the treatment of cancer, interaction of the cancer with the environment as well as for looking at drugs, pharmaceuticals and other chemical interactions. The transgenic animals are useful in any assay in which the skin cells of the animal can be used.

One specific embodiment of the present invention is a method for the enhanced healing of a wound or surgical incision. This method comprises the *in vivo* transduction of epidermal cells with a keratin K1 vector. The nucleic acid cassette of said vector contains a nucleic acid sequence for a growth factor.

In the preferred embodiment for the treatment of wounds or -surgical incisions, a plurality of vectors are introduced into the epidermal In the plurality of vectors, the cassette of at least one vector contains a nucleic acid sequence for an epidermal growth factor (TGF-a), the cassette of at least one vector contains a dermal growth factor (PDGF), a cassette of at least one vector contains a nucleic acid sequence for a matrix protein to anchor the epidermis to the dermis, and a cassette of at least one vector contains a nucleic acid sequence for an angiogenesis The sequence for matrix proteins can be selected from any sequences useful for the anchoring of the epidermis to the dermis but are usually selected from the group consisting of Type IV collagen, laminin, nidogen, and Type VII collagen. The angiogenesis factor is usually selected from the group consisting of acid fibroblast and basic fibroblast growth factors, and angiogenin. The combination of the vectors provides all of the necessary elements for quick and rapid enhancement of healing of wounds or surgical incisions. This procedure is very helpful in the case of plastic or reconstructive surgery. Furthermore, skin ulcers can be

treated by following similar procedures as described for wound healing or surgical incision. These procedures for healing of wounds, surgical incisions and skin ulcers are useful in animals and humans.

In the ex vivo approach for treating or healing wounds, surgical incisions and skin lesions, the vectors are first transduced into the epidermal cells ex vivo. The transformed epidermal cells are transplanted onto the animal or human to be treated.

Another embodiment of the present invention is a method for treating psoriasis. In this method, epidermal cells are transduced in *in vivo* with a keratin K1 vector. A nucleic acid cassette in said vector contains a nucleic acid sequence for a protein or polypeptide selected from the group consisting of TGF-β, a soluble form of cytokine receptor, and an antisense RNA. The cytokine receptor can be selected from the group consisting of IL-1, IL-6, and IL-8. The antisense RNA sequence is selected from the group consisting of TGF-α, IL-1, IL-6, and IL-8.

In another embodiment of the present invention there is a method of treating skin cancer. This method comprises the steps of *in vivo* transduction of epidermal cells with a keratin K1 vector. The nucleic acid cassette of either vector contains the nucleic acid sequence coding for antisense RNA for the E6 or E7 genes of the human papilloma virus or coding for the normal p53 protein.

It has been found that the keratin K1 vector contains a novel negative regulatory element in its 3' flanking sequence which can be suppressed by Vitamin D₃. With the Vitamin D₃ regulatory element in the vector, the expression of a nucleic acid cassette can be regulated by Vitamin D, a commonly used substance in animals and humans.

The human keratin K1 vector can also be modified by the insertion of additional 5' flanking sequences from an 18 kb Eco RV fragment to its 5' end (nucleotides 6090 to 14180 of SEQ. ID. No. 3). The addition of these sequences allows the human keratin K1 vector to be expressed

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exactly like the endogenous K1 gene, that is, post mitotically in cells committed to terminal differentiation. Since these cells are programmed to die and will eventually slough into the environment, this is another way of producing transient expression in cells.

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An additional embodiment of the present invention is a method for vaccination comprising the step of *in vivo* introduction of a keratin K1 vector into epidermal cells. The nucleic acid cassette in the vectors usually codes for a polypeptide which induces an immunological response. An example of this is the viral capsid from the human papilloma virus. One skilled in the art will readily recognize that any other variety of proteins can be used to generate a immunologic response and thus produce antibodies for vaccination.

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The following examples are offered by way of illustration and are not intended to limit the invention in any manner.

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EXAMPLE 1

Construction and Characterization of a Vector From the HK1 Gene To target the expression of exogenous DNA to the epidermis a vector from the human keratin K1 gene was constructed. Among its many uses, it is useful in making transgenic animals.

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A schematic showing the structure of the human keratin K1 gene is shown in Figure 1. The 12 kb EcoRI fragment containing the entire human keratin K1 gene was originally isolated from lambda clone c55 (Johnson, et al., PNAS, USA, Vol. 82, pp. 1896-1900, (1985)). In constructing the targeting vector, most of the first exon including the ATG was removed, leaving only the 5' non-coding sequences, the first intron and the intron-exon boundaries. In addition, the remainder of the gene up to the termination codon was deleted. A poly linker containing the following unique restriction sites (Bam HI, Xma I, Kpn I, Not I, and Cla I) was engineered into a site 3' of the first intron to allow easy insertion

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of exogenous DNA. These manipulations were performed through the use of polymerase chain reactions (PCR). The unique EcoRI sites were conserved at the ends of the vector to allow easy amplification in pGEM vectors and excision for purification from plasmid sequences prior to injection into embryos.

The rationale for constructing the vector in this manner was as follows. Since the specific elements responsible for the expression characteristics of the 12 kb human keratin K1 fragment have not been defined, the entire 5' and 3' flanking regions were included in the vector construct. One skilled in the art will readily recognize that as these elements are further defined the flanking sequences can be changed accordingly. In addition, sequences within the 3' non-coding region were retained since these may confirm stability to transcripts of exogenous DNA in epidermal cells. The first intron was retained to potentially enhance expression efficiency (Brinster, et al., PNAS, USA, Vol. 82, pp. 1896-1900 (1988).

EXAMPLE 2

HK1 Expression in Epidermal Keratinocytes

To assess the human keratin K1 targeting vector for exclusive expression in epidermal keratinocytes, the β-galactosidase reporter gene was cloned into Bam HI and Cla I restriction sites located in the polylinker region of the expression vector (Figure 1). The β-galactosidase gene has frequently been used as a reporter gene to assess targeting specificity (MacGregor et al., In: Methods in Molecular Biology Vol. 7, pp. 217-235 (1991). This construct was designated pHK1.β-gal. To determine if expression of this construct resulted in the production of a functional protein, and to determine whether the vector retained cell type specificity, this construct was transfected into primary epidermal keratinocytes and primary dermal fibroblasts. At seventy-two hours post transfection cells

were stained with a solution containing the substrate 5-bromo-4-chloro-3-indoyl- β -galactosidase (X-gal). β -galactosidase activity, indicated by a blue coloration, was detected in keratinocytes but not fibroblasts. Thus, expression of the HK1. β -gal construct was cell type specific and resulted in the production of a functional protein.

EXAMPLE 3

Transgene Mice

The same pHK1. β -gal construct utilized in the *in vitro* studies discussed in Example 2 was used in the production of transgenic mice. This construct was digested with EcoRI (see Figure 1) and subjected to preparative agarose gel electrophoresis to purify the pHK1. β -gal expression construct away from plasmid sequences (pGEM 3) which might interfere with expression. The separated expression construct sequences were purified and recovered using NA 45 DEAE membrane (Schleicher & Schuell). DNA was precipitated and resuspended at 1-3 ng/ul. ICR outbred female mice (Sasco) were given PMS and HCG to stimulate superovulation, mated to FVB males (Taconic) and the resulting early fertilized embryos (most preferably on cell stage) were collected from the oviducts. DNA was micro-injected into the pronuclei and the embryos were surgically transferred to pseudopregnant recipient females (the result of mating ICR females with vasectomized $B_6D_2F_1$ males (Taconic))

In the initial experiments, 40 mice were born. In order to quickly determine if the pHK1.β-gal transgene was being exclusively expressed in the epidermis of these mice, these animals were sacrificed at birth. A small amount of tissue was removed for extraction of DNA and the remainder of the neonate was rapidly frozen in Tissue-Tek O.C.T. for frozen sections. PCR analysis was performed on the extracted DNA using oligonucleotide primers specific for the intron within the HK1 vector and

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this demonstrated that 5 of the 40 neonates contained the HK1.β-gal construct.

To assess whether expression of the HK1β-gal construct was restricted to the epidermis or expressed in other squamous epithelia, frozen longitudinal sections were cut from several PCR positive and PCR negative embedded neonates and these were stained with X-Gal. Typical results are shown in Figure 2 where PCR positive animal, #30, expressed high levels of β-galactosidase in the epidermis (Figure 2A and a PCR negative sibling, #29, was completely negative (Figure 2B), indicating that endogenous murine B-galactosidase was not expressed at sufficient levels in the epidermis to cause false positives in this assay. Staining of the intestine was observed in both the positive (#30) and negative (#29) This may represent endogenous enzyme activity or the neonates. production of β-galactosidase by bacteria in the intestine. X-gal staining was detected in the basal compartment, although it is not as intense as in the differentiated layers (Figure 2D). Thus, the human keratin K1 expression vector is also expressed in a substantial number of proliferating basal cells.

The most important finding from these initial transgenic experiments in that the vector constructed from the human keratin K1 gene can target the expression of an exogenous coding sequence exclusively to the epidermis of transgenic mice. This specificity of targeting can be readily seen in Figure 2A. This low power exposure of the skin of #30 demonstrates intense staining with X-Gal. In addition, there are numerous hair follicles and sebaceous glands in this section which are marked by arrows and these do not stain with X-Gal. Keratins K5 and K14 are not only expressed in the epidermis, but in all squamous epithelia, including hair follicles and sebaceous glands. The expression pattern for keratin K14 (Figure 2C) is revealed by immunofluorescence with a specific K14 antiserum of an area from a consecutive section that

is comparable to that in Figure 2A. Note staining of the epidermis, as well as, hair follicles and sebaceous glands. If the strategy used in construction the human keratin K1 expression vector had altered its targeting specificity in transgenic mice, then X-Gal staining would have been observed in hair follicles, sebaceous glands, other squamous epithelia, and perhaps even other tissue types. However, expression of the HK1. β -gal transgene, like the keratin K1 gene itself is restricted to the epidermis.

EXAMPLE 4

Regulation of Keratin K1 Vector by Vitamin D₃

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A novel Vitamin D_s responsive element was used to modulate expression levels in the epidermis. Although all of the regulatory elements of the human keratin K1 gene have not been identified, a novel negative regulatory element from the human keratin K1 gene (HK1.NRE) has been identified and this example demonstrates that it is able to suppress a heterologous promoter in response to Vitamin D_s. The HK1.NRE is 70 nucleotides in length (nucleotides 9134 to 9204 of SEQ. ID. No. 1). PCR technology was used to generate Bam HI and Bgl II sites at opposite ends of this fragment. This facilitates generating multiple copies of this fragment since ligation and digestion with Bam HI and Bgl II will select for oligomers which have ligated head to tail. Four tandem copies of the HK1.NRE were inserted into the Bgl II cloning site of pA10.CAT. In the absence of Vitamin D, this construct is highly expressed when transfected into primary mouse epidermal cells (Figure 3). The addition of increasing concentrations of Vitamin D₃ to the culture medium completely suppresses transcription of this heterologous promoter. This observation indicates that the activity of the human keratin K1 expression vector can be modulated in the epidermis. The activity of the human keratin K1 vector is suppressed in the epidermis by topical application of Vitamin D_s, or an analogue, to the skin.

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EXAMPLE 5

Development of Transgenic Animal Models for Skin Carcinogenesis

The ability to stably introduce genes into the germline of mice has greatly enhanced prospects for generation of animal models of human disease (Leder and Stewart U.S. Patent No. 4,736,866 issued April 12, 1988 and Palmiter and Brinster, Ann. Rev. Genet., Vol. 20, pp. 465-499). When such genes are combined with regulatory sequences that target their expression to specific tissues, it provides a model to not only study diseases in the context of living organisms, but also in specific tissues suspected of being the targets of these genes. Thus, transgenic mice offer the possibility to determine the influence of factors such as blood supply. an intact immune system, humoral and cell-mediated growth controls and physical barriers on disease progression. The epidermis is an attractive tissue for targeted gene expression; not only is it a model for epithelial diseases in general but the accessibility of the epidermis allows easy detection of progressive pathological changes that result from transgene expression as well as the assessment of the potential role played by environmental factors in these processes. In addition, the prospects for utilizing gene therapy to treat cancer are coming closer to reality. Therefore, animal models of human cancers would be useful to assess the therapeutic potential of these approaches. The development of animal models of skin disease is dependent upon the ability to specifically target gene expression to the epidermis. The human keratin K1 targeting vector described in Example 1 is ideally suited for this purpose.

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EXAMPLE 6

Targeting the v-ras^{Ha} Oncogene to the Epidermis

One family of proto-oncogenes, the ras family (ras^{Ha}, ras^{K1}, ras^N) has been identified in approximately 20% of human tumors by virtue of specific point mutations at codons 12, 13, and 61 which activate their

transforming potential. The mechanisms whereby ras genes become activated are currently unknown but there is widespread evidence that environmental agents play pivotal roles in the etiology of ras mutations. To date few studies have undertaken to study ras activation in human skin malignancies. However recent reports have identified ras Ha activation in basal and squamous cell carcinomas appearing on sun exposed body sites, interestingly at potential pyrimidine dimer sites possibly derived from skin exposure to UV irradiation. In the mouse skin model of chemical carcinogenesis where the three distinct stages of initiation, promotion and malignant conversion have been defined, ras Ha activation has been found in benign squamous papillomas, the end point of initiation and promotion suggesting an early role for ras^{Ha} in skin carcinogenesis. Taken collectively, the above experimental evidence suggests the importance of developing an animal model to further study the mechanism of ras Ha-induced skin carcinogenesis. Toward this end, the sequence encoding the v-ras^{Ha} protein of Harvey Murine Sarcoma Virus (Dhar, et al., Science, Vol. 217, pp. 934-937, (1982) was cloned into the Bam HI and Cla I sites of the human keratin K1 expression vector (Figure 4) discriminate expression of the v-ras Ha transgene from that of the endogenous ras gene, a sequence encoding the human keratin K6 epitope SEQ. ID. No 4 was engineered onto the 5' end of the v-ras^{Ha} cassette.

HK1 ras transgenic mice exhibit the following phenotype: 1) Newborn transgenic mice expressing v-ras^{Ha} (HK1 ras) exclusively in the epidermis show distinct wrinkled skin at 48 hours and are smaller than litter mates. 2) Juvenile HK1 ras transgenic mice exhibit progressive keratinization which peaks at 14 days. 3) The histotype of newborn HK1 ras mice reveals massive epidermal hyperplasia with up to 20 fold thickening of the epidermis. 4) By day 14 this progresses to massive hyperkeratosis. Both histotypes are pre-neoplastic, papillomatous, non dysplastic and exhibit few appendages.

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The HK1 ras transgenic mice develop benign tumors. Typical lesions appear within 10-12 weeks at single sites. The histotype of these tumors reveals a well differentiated squamous papilloma. Papillomas often appear at sites after wounding. Many of these papillomas are prone to regression. This regression phenomenon suggests that ras^{Ha} alone is insufficient to maintain even a benign phenotype and requires further events which may involve roles for additional oncogenes/antioncogenes.

EXAMPLE 7

Targeting the fos Oncogene to the Epidermis of Transgenic Mice

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Recent in vitro studies have shown that the v-fos gene can convert to malignancy primary keratinocytes or papilloma cell lines which expressed an activated ras^{Ha} (Greenhalgh, et al., PNAS, USA, Vol. 87, pp. 643-647, (1990); Greenhalgh and Yuspa, Mol. Carcinogen., Vol. 1, pp. 134-143 (1988). This suggested that fos could play a later role in epidermal carcinogenesis and cooperate with the benign phenotype imparted by activated ras-R expression. Although this alone was sufficient to initiate the establishment of HK1 fos transgenic mice with a view to mate with HK1 ras mice, two further studies have identified a role for fos in normal epidermal differentiation and thus highlights fos as an attractive target for perturbation. Using a c-fos/β-gal fusion gene Curran and co-workers (Smeyne et al, Neuron, Vol. 8, pp. 13-23 (1992)) have shown significant fos expression in the differentiated layers of the epidermis and (Fisher, et al., Development, Vol. III, pp. 253-258, (1991)) have localized c-fos expression to a specific subset of granular cells. Thus fos may have an important role in the control of the final stages of keratinocyte differentiation. The putative perturbations of this normal role for c-fos in such specialized cells by v-fos can only be explored in the context of targeted expression in transgenic mice. In addition the c-fos proto-oncogene is known to function as a transcriptional regulator in conjunction with the c-jun/AP1 gene

product and thus, while targeting ras^{Ha} represents studies of membrane signalling on neoplasia, targeting fos explores the role of transcriptional control on this process.

Thus, the fos protein coding sequence from the FBJ/FBR chimeric v-fos plasmid pFBRJ was inserted into the human keratin K1 targeting vector (Figure 5). To discriminate expression of the v-fos transgene from that of the endogenous fos gene, a sequence encoding the human keratin K1 epitope (SEQ. ID. No. 5) was engineered onto the 5' end of the v-fos cassette.

HK1 fos transgenic mice exhibit the following phenotype: 1) A specific ear phenotype typically appears at 3-4 months initially in the wounded (tagged) ear and then becomes bilateral. 2) In several animals expressing severe phenotypes, the wounded ear lesion can grossly resemble a benign keratoacanthoma. 3) Alopecia and hyperkeratosis of the axilla often develop in older animals (approximately 1 year of age).

The histotypes of the HK1 fos mice are as follows: 1) The histotype of the initial ear lesions exhibits hyperplasia and hyperkeratosis, a pre-neoplastic pathology with few dysplastic cells and little evidence of further neoplastic progression. 2) At later stages the massive hyperkeratotic histotype resembles a benign keratoacanthoma.

Three HK1 fos transgenic mice lines have been established which develop an obvious pre-neoplastic ear phenotype at 3-4 months. The promotion stimulus derived from wounding (i.e. ear tag) appears to accelerate the appearance of this phenotype which eventually becomes bilateral. Also, it appears that friction in the axilla and inguinal area may also promote a pre-neoplastic hyperplastic/hyperkeratotic response after a significant latent period. Collectively these data support a fundamental role for the fos gene in normal keratinocyte differentiation and perturbation by v-fos results in pre-neoplastic differentiation disorders. In several HK1 fos mice severe ear lesions appear to progress to resemble

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benign keratoacanthomas. Although numbers are low at this time, that this is the resultant tumor type is consistent with a role for fos in the latter stages of terminal differentiation, and low numbers and latency suggest a requirement for additional events.

EXAMPLE 8

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Targeting HPV 18 E6 and E7 Gene Expression to the Epidermis

There is widespread evidence from clinical and epidemiological studies which implicate human papilloma viruses (HPV) in the etiology of certain squamous epithelial tumors in humans. HPV's have a specific tropism for squamous epithelial cells and different types of HPVs have specificity for the anatomic site that they infect. Additionally, within a specific subgroup of HPVs, certain types are associated with development of either benign (e.g. HPV6 and 11) or malignant (e.g., HPV-16 and 18) disease and this may center on the properties of the E6 and E7 genes. Through adaptation to the differentiation programs of the epithelia that they infect, HPVs have evolved a clever strategy for the production of infectious progeny. HPVs infect basal epithelial cells but do not undergo lytic replication in this compartment, thus, the germinative pool of cells is not subjected to the cytopathic effects of late viral gene expression. Production of virus only occurs in terminally differentiated cells that have lost proliferative potential and will be desquamated into the environment. This strategy not only provides for the spread of mature viral particles, but ensures their continuous production by replenishment with cells from the basal compartment. Since the life cycle of the virus is so tightly linked to all stages of differentiation of squamous epithelial cells, establishment of successful culture systems has been difficult. To date, these host factors, coupled with regulatory mechanisms present within papilloma virus genomes themselves have also hindered attempts to observe pathological effects of HPV gene expression in squamous epithelia in

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transgenic mice. These restrictions on utilization of the transgenic mouse model have been overcome with the ability to specifically target HPV gene expression to squamous epithelia using the human keratin K1 targeting vector. In the example provided, the coding sequence for the E6 and E7 genes of HPV 18 were inserted into the human keratin K1 targeting vector at the Bgl II and Cla I sites (Figure 6).

HK1 E6/E7 mice exhibit the following phenotypes and histotypes:

1) One mouse exhibited a subtle skin lesion at 7 months characterized by skin rigidity, thickening and roughness underlying the fur which later progresses to a wart like structure by 10 months. 2) The histotype of this lesion exhibits hyperplasia, hyperkeratosis and the beginnings of verrucous formation. 3) The histotype of a lesion from another mouse at 11 months, is characteristic of a typical wart induced by HPV.

To date three HK1 E6/E7 transgenic mouse lines have been established which develop HPV-like lesions with low frequency and long latent periods. At this time it is unclear whether this limited appearance of phenotypes reflects the subtle nature of the lesion, and its requirement for a long latency period, or the complex nature of HPV biology. However it is noteworthy that our result is consistent with the epidemiology of HPV infections in humans, e.g. although a large percentage of a given female population can test positive for cervical infection by HPV 6, 11, 16, and 18 relatively few progress to develop overt lesions. It may be therefore that the apparent delay and low phenotype frequency exhibited by these mice provides a relevant background to study the consequences of HPV expression during epithelial differentiation. In addition, these mice can be useful in assessing the efficacy of novel antisense pharmaceuticals which have been designed to inhibit expression of the E6 and E7 genes of HPV 18.

EXAMPLE 9

Production of Transgenic Mice Expressing TGF- α in the epidermis

Transforming growth factor alpha (TGF- α) is a cytokine with structural and functional characteristics similar to epidermal growth factor (EGF). Both TGF- α and EGF bind to the epidermal growth factor receptor (EGF-R) and stimulate the tyrosine kinase cascade. TGF- α is expressed by both normal and transformed cells and causes proliferation of cultured keratinocytes. In vivo, TGF- α induces angiogenesis and is more potent than EGF in accelerating wound healing. In normal human skin, expression of TGF- α occurs in all layers of the epidermis and in certain areas of the appendages. Several cutaneous diseases such as psoriasis, squamous cell carcinoma, and congenital bullous ichthyosiform erythroderma have been associated with altered expression of TGF- α .

To determine whether altered expression of TGF- α plays a role in the pathogenesis of these diseases, the protein coding sequence of human TGF- α was inserted into the human keratin K1 targeting vector (Figure 7). Injection of the HK1.TGF- α construct into embryos resulted in phenotypic founders that were quite similar to that of ras^{Ha} . The histotype was also similar with epidermal hyperplasia, hyperkeratosis and relative alopecia. One founder (2 1/2 months of age) has developed multiple papillomas. Histologically these appear to be squamous papillomas. To date, none of these lesions have converted to a malignant phenotype.

EXAMPLE 10

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Production of Transgenic Mice Expressing the HIV tat gene in the epidermis.

Patients infected with the human immunodeficiency virus (HIV) are at high risk for the development of specific AIDS-associated cutaneous disorders. Often patients manifesting symptoms have skin lesions ranging

from hyperproliferative conditions such as psoriasis to Kaposi's sarcoma and metastatic basal cell carcinoma. The precise role of HIV genes, the cells of origin and hence etiology of such skin lesions remains unknown. It may be that specific HIV genes, e.g., the trans-regulatory protein tat, play a role directly or indirectly on the homeostatic mechanisms of host cells and tissues. Alternatively, the HIV tat gene may interact with or activate other viral genes present from latent or opportunistic infections, e.g., human papilloma virus (HPV). To directly assess the role of keratinocytes in the development of AIDS-associated cutaneous disorders, the HIV tat gene is targeted to the epidermis of transgenic mice. Targeting of the tat gene and exclusive expression in keratinocytes is achieved by the use of the human keratin K1 vector (Figure 8). The development of strains of mice which develop cutaneous lesions with predictable kinetics as a result of expression of the HIV tat gene alone or in combination with other oncogenes serves as a useful model for assessing therapeutic potential of antisense pharmaceuticals designed to inhibit expression of the HIV tat gene.

EXAMPLE 11

Utilization of the HK1 Vector for Gene Therapy Applications

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Where exclusive expression in epidermal cells is desirable and for transient expression the HK1 vector is an excellent choice for gene therapy. Unlike the human keratin K1 gene itself, the human keratin K1 vector derived from the 12 kb fragment is expressed in proliferating basal cells in the epidermis. In more recent transgenic experiments, it has been determined that a larger fragment containing the human keratin K1 gene, a 18 kb Eco RV fragment (Shown schematically in Figure 9), is expressed exactly like the endogenous mouse K1 gene, i.e. post mitotically in cells committed to terminal differentiation. These cells are programmed to die and will eventually slough into the environment. Therefore, for human

applications where transient expression is desired, it is possible to design a vector that will only be expressed in cells after they commit to terminal differentiation and begin moving upward toward the outer layers of the epidermis. The vector will be expressed approximately 10-14 days prior to being shed into the environment. This can be accomplished by inserting additional 5' flanking sequences from the 18 kb Eco RV fragment onto the end of the original human keratin K1 vector (See Fig. 10).

EXAMPLE 12

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Detection of Carcinogens and Tumor Promoters

Short-term tests (STTs) for genotoxic chemicals were originally developed as fast, inexpensive assays to assess the potential hazard of chemicals to humans. However, a recent report summarizing the results of a project initiated by the National Toxicology Program to evaluate the ability of STT's to predict rodent carcinogenicity questions the validity of relying solely on STT's. Three of the most potent carcinogens, detected in the rodent assays, produced no genetic toxicity in any of the four STTs evaluated (Tennant et al., Science, Vol. 236, pp. 933-941, (1987). Thus, to receive EPA/FDA approval for new compounds, chemical, agricultural, food and drug companies are currently required to perform two year animal tests costing up to \$2 million. The development of new transgenic strains of mice that have been genetically engineered to rapidly detect carcinogens and tumor promoters would substantially reduce the overhead cost of long-term animal studies. The suitability of the transgenic mouse lines claimed in this patent application for rapid detection of carcinogens is initially determined with a known skin carcinogen, DMBA, to determine whether benign lesions appear earlier than in control non-treated litter mates. To determine suitability for detecting tumor promoters, a known promoter, 12-O-tetra-decanoylphorbol-13-acetate (TPA) is applied to ras^{Ha} ,

and fos mice. Since benign lesions in ras^{Ha} mice and hyperplasia in fos mice appeared at sites of wounding (i.e., tagged ears), and wounding can promote tumor formation, these lines are useful for these studies.

All patents and publications mentioned in this specification are indicative of the levels of those skilled in the art to which the invention pertains. All patents and publications which are incorporated herein by reference are incorporated to the same extent as if each individual publication was specifically and individually indicated to be incorporated by reference.

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One skilled in the art will readily appreciate that the present invention is well adapted to carry out the objects and obtain the ends and advantages mentioned, as well as those inherent therein. The bioreactors, nucleic acid sequences, transformed epidermal cells, transgenic animals and human keratin K1 vector, along with the methods, procedures, treatments, molecules of specific compounds, are exemplary and are not intended as limitations on the scope of the invention. Changes therein and other uses will occur to those skilled in the art which are encompassed within the spirit of the invention as defined by the scope of the claims.

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SEQUENCE LISTING

(i) APPLICANT: Roop, Dennis R.
Rothnagel, Joseph A.
Greenhalgh, David A.
Yuspa, Stuart H.

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- (iii) NUMBER OF SEQUENCES: 5

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- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Fulbright & Jaworski
 - (B) STREET: 1301 McKinney, Suite 5100
 - (C) CITY: Houston
 - (D) STATE: Texas
 - (E) COUNTRY: U.S.A.
 - (F) ZIP: 77010-3095
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
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 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
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 - (A) NAME: Paul, Thomas D.
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(ix) TELECOMMUNICATION INFORMATION:

	(A) TELEPHONE: 713/651-5325	
_	(B) TELEFAX: 713/651-5246	
5.	(C) TELEX: 762829	
5	(2) INFORMATION FOR SEQ ID NO:1:	
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 10747 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: double	
10	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: DNA (genomic)	
	(iii) HYPOTHETICAL: NO	
	(iv) ANTI-SENSE: NO	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
	(x2) Dagomica Substitutions dag do most	
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		420
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		540
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35	TGGGAGCTGC	TGCAGCAGGT	AGATACCTCC	ACTAGAACCC	ATAATTTAGA	GCCCTACTTT	3000
	GAGTCATTCA	TCAACAATCT	CCGAAGGAGA	GTGGACCAAC	tgaagagtga	TCAATCTCGG	3060
	TTGGATTCGG	aactgaagaa	CATGCAGGAC	ATGGTGGAGG	ATTACCGGAA	CAAGTAAGGG	3120
	ACCCTGTCTG	GGCAGTTCTT	AACTTTTGCT	GTAAAAGAGT	TCCAGAAAGT	aataagctaa	3180
	CATCATGAAG	CACCATGTAG	CTATGTCTTT	TCTTAGGTTA	GAGGCACATC	AGTTTGACAT	3240

:	TTTCAGAAAT CTTCATTTTC TCAGGAGATG GAAATAGTCT AGTGGTTTTA TTGCTCAGTA 33	DO
	GAAAGTAGTG GCCAATATGT CCTAGGTTCA TAATAGAAAG GCAGTGATAG GCAATGCCAC 33	60
	CTTTAGTTTA GAATGCTGGA CTTCAGGTCT TACCACCTCT GAATCTCCTA ATTGTTTCTG 34	20
ie.	CTTTCCTGCA GGTATGAGGA TGAAATCAAC AAGCGGACAA ATGCAGAGAA TGAATTTGTG 34	80
5	ACCATCAAGA AGGTAAGCAA ATTCTGTAGG ACGGAACTCA CATTTGAAAT AAATAAGGGA 35	40
	AGAGGGTCTC CAATTACTAA GCAGAAAGCA GCCATGATAT GGAGAGCCAG GTAGTAGACC 360	20
	TGGGGAGTAT ATGGAGTGGG GCTATATTTT TCACATCATC ATGGACCTGG ACTGATCCAG 36	50
	GCACTTGGCT TCTCCATATT TCCCAGCACC TTACATAGTA AGTGGAGTGG	20
	GCAAGCCAGG CACACTCCCT TGATGGTGCT ATCCGGGGGT GGGACAGTTA GGGAACTGTG 378	30
10	ATTTACCTGG GGCAAAAAGG AGTGGAGTAG ACCCAAAGCT CCTTTTTTTG CTTGGAGAAT 384	40
	CCCCTCACAG GTAATGAGAG GGACCTGCCC TGGAGAGAAC GTGCCTTCAT GATGTCCCTT 390	00
	GTTCCTCTAG GATGTGGATG GTGCTTATAT GACCAAGGTG GACCTTCAGG CCAAACTTGA 396	50
	CAACTIGCAG CAGGAAATIG ATTICCTIAC AGCACTCTAC CAAGCAGTAA GICTICCAGI 402	20
	TTCAACCAAG TTTATCTAAA TGGAGAGTTT TTAAGCCGGA ACCCACAACG ATTCAGAAGA 408	30
15	ATAGATATTT ATCTTTATT TCCTGACTGC TTTCTCTGTC TAAGTTGTTT TTTGTTTTAG 414	10
	TGCTGTAAGA GTCACTAACC TATTATGTCT TGCAGGAGTT GTCTCAGATG CAGACTCAAA 420	00
	TCAGTGAAAC TAATGTCATC CTCTCTATGG ACAACAACCG CAGTCTCGAC CTGGACAGCA 426	50
.	TCATTGCTGA GGTCAAGGCC CAGTACGAGG ATATAGCCCA GAAGAGCAAA GCTGAGGCCG 432	20
	AGTCCTTGTA CCAGAGCAAG GTGAGTGGGC TGAAACCGTA GCCAGTTTCC CTGAAATGGC 438	30
20	TTGTCTTGCT ATCCTGTGTT ATCTCATGTA TGTGTGCCTG TGCCATGCTG AGTTCTGCCT 444	10
	ACATTTAACA AACGCTATCT ACCATCTTTA GTATGAAGAG CTGCAGATCA CTGCTGGCAG 450)0
	ACATGGGGAT AGTGTGAGAA ATTCAAAGAT AGAAATTTCT GAGCTGAATC GTGTGATCCA 456	50
	GAGACTTAGA TCTGAAATCG ACAATGTCAA GAAGCAGGTA TGTGCTTTCT CCTTCTACCA 462	20
	CTCAGCTGTA TGGAATGGGG GTAACCCTCA GGTAAAGGGC GAGTGCTTTC CTAGTTTTGA 468	30
25	ATCTTGCAAT TCAGCCCAAG GCTACATTAT TAGCCCTGGT TCCTTTTCTG ACTATGCTAG 474	10
	TTTCCAGAAT GCAGCCATCA TGCTGGGTTC TCTTTAGGGA AATCTGTGAG AATGGCCTAG 480	00
	TAGAGAAAGA TGGGATGGTC AATGTGAGTG ATCTAGCCTA TGACCCAAAG TGGACTTAAG 486	50
	AGTTGGGGAG TGAGAGGAAG GGCAGCCAGG AGGTTTTAGA GTAGGTGTTT AGAAGAATGT 492	10
	CAAGTCTGTA AGGGTTGTAG GAGCCTTGAC TCAGGGCCAA GAGAGGCTGT TGAGTTATCC 498	30
30	CTANGGTCTT TTANGGANGT CANCATGGTG ATGTGTTATC TGGAGGTGGG TGTGAGATGA 504	10
	CTTAAGGCCA AGTGGTTCTG TTGGACTCAT TATTGGCCTC ACTGGAGTGG GGAGACCAAT 510)0
	TGGGATGAGG AGGCCTAGTG GGGAATGCAT ATTATGAGAG GGTGTCATAT CTTTTTCAGA 516	
	TCTCCAACTT GCAGCAGTCC ATCAGTGATG CAGAGCAGCG TGGCGAGAAT GCCCTCAAGG 522	! O
	ATGCCAAGAA CAAGCTGAAT GACCTGGAGG ATGCCCTGCA GCAGGCCAAG GAAGACCTGG 528	10
· 35	CCCGCCTGCT GCGTGACTAC CAGGAGCTGA TGAACACCAA GCTGGCCCTG GATCTGGAGA 534	10
	TTGCCACCTA CAGGACCCTC CTGGAGGGAG AAGAAAGCAG GTGAGGAAGG GACGCTGGGA 540	10
,	GTCGAACCTC TTCTCATGGT CTTCCTTCCT TGCAAGCTGA TTGTTGTTGA AGATGCAGCC 546	0
	ATCTGATTGC AGCTTGTGCT GGGTATGGGG AAATGAAAAG TACACGGAGC AGGAGGAAGG 552	: O
	ACCTAGTTTT ACTTTGGGAG CTGGAGTCCC AAGCTGTTTA TTTTTTTCTT CTAGGGCTGT 558	10

	AACATATCTA GAAAGAGCTT TGAGGTGGAG CAAATTATTC TTTATCTGGG CTGCCTCAGA	5640
	TGGCAGCTGG CCTAAAGTCG GCATCTTTAG AGGGGGCCTT CATTGGCTGC AAGGCTCGTC	5700
	TCGTTTATAT GGGAATTTCT CCGTGTTTGT ACTCTTGCTG AGAAAAAATG ACAGGTCTGG	5760
	GAGGCCAGAG GGGATTGGAT TAAGTTTCAG ATTAAGTGCA TTGGAGAAGA CCCAGATGGG	5820
5	GAAAGTCTTC AAGGTGGTGG AGCGGGGAAT GGGGAAGCGG TITGGGAAGC TGGAGTGTCC	5880
	TGAGGAATTT TCTTATTTTC TCCTACAGGA TGTCTGGAGA ATGTGCCCCG AACGTGAGTG	5940
-	TGTGTAAGTA CAAGTCGATT TCTCAGGGGC ATGTGCAGGC TTTGTTGGGC TGGAAACGGA	6000
	GTTGAGGTTG AAAATAACTG AGCTTCCTCT TGCAGCTGTG AGCACAAGCC ACACCACCAT	6060
	CAGTGGAGGT GGCAGCCGAG GAGGTGGCGG CGGTGGCTAC GGCTCTGGAG GTAGCAGCTA	6120
10	TGGCTCCGGA GGTGGTAGCT ATGGTTCTGG AGGTGGCGGC GGCGGCGGCC GTGGCAGCTA	6180
	TGGCTCCGGA GGTGGCAGCT ATGGCTCTGG AGGTGGCGGC GGCGGCCATG GCAGCTACGG	6240 ·
	CTCCGGAAGC AGCAGTGGGG GCTACAGAGG TGGCTCTGGA GGCGGCGGCG GCGGCAGCTC	6300
	TGGCGGCCGG GGCTCTGGCG GCGGGAGCTC TGGAGGCTCC ATAGGAGGCC GGGGATCCAG	6360
	CTCTGGGGGT GTCAAGTCCT CTGGTGGCAG TTCCAGCGTG AAGTTTGTTT CTACCACTTA	6420
15	TICCGGAGTA ACCAGATAAA GAGATGCCCT CTGTTTCATT AGCTCTAGTT CTCCCCCAGC	6480
	ATCACTARCA ARTATGCTTG GCAAGACCGA GGTCGATTTG TCCCAGCCTT ACCGGAGAAA	6540
	AGAGCTATGG TTAGTTACAC TAGCTCATCC TATTCCCCCA GCTCTTTCTT TTCTGCTGTT	6600
	TCCCAATGAA GTTTTCAGAT CAGTGGCAAT CTCAGTCCCC TGGCTATGAC CCTGCTTTGT	6660
	TCTTTCCCTG AGAAACAGTT CAGCAGTGAC CACCACCCAC ATGACATTTC AAAGCACCTC	6720
20	CTTAAGCCAG CCAGAGTAGG ACCAGTTAGA CCCAGGGTGT GGACAGCTCC TTAGCATCTT	6780
	ATCTCTGTGC TGTTTTGGTT TTGTACATAA GGTGTAAGCA AGTTGTTTTT CTTTTGTGGA	6840
	GAGGICITAA ACTCCCCATT TCCTTGTTTT GCTGCAATAA ACTGCATTTG AAATTCTCCA	6900
	TGTCTCGATC GCCCTTGTTT ACGGCACTGT CTAACCTGGA TGGGTGTTTT GTGAGGTAAA	6960
	AGAAGACACT AGAGCCACAT GGCATATGGG AAAGTCATGC ACACAAACAT GAGAAAAATG	7020
25	CAGAGGCCAA CCAGGCAACA TTTCACCAGA CTGGAATCAC AGAGAGAGCA AGCACTTTCC	7080
	CAGATGGTGG GGATGTCATG GAGAAATGGA GAGACCGGGT GACAGGTTTT GTTCATTTGA	7140
	GAAGGCTTTC TTGAAAAGGG CAGTGAGCAA GCAGGTTGGG AGGAAGAGGT GTGGCATTGA	7200
	GAAGAAGGGA AAGTATTGCA TGAAAAAGTA ATTCTTCACG TGGAACAGCC AGTAAGGAGG	7260
	GGCATGAGTA ATATAGGGTC AGCAGTTACT GGAGCCAGAA TACAGACTTT GGCCTGGGGA	7320
30	GTTCAAGAAC TAAGAGTGGT AATAGAGAGT TGGATATTCC ATTTCCCTTC TCTTTTTGTG	7380
	CCACCACCCA AAGCTCTGCA TAATCTAAGA AGTTCCCTTG TTGACACATA GCTCATACTT	7440
	GTGAAGTTGT ACAACAGGAT AGCATAGTGG CCAGAAGCAT GGACAGTTGA ACTCAGATAT	7500
	GCTTGGGTTT GAATCTTACC ATCACCATTT ACTAGTTCTG TAATACAGTG CAAGTTACAG	7560
	ACATCTCTGC ACCTCAGTTT TAGTATGTCT AAATTGGGGA TGATAATGCC TTCCTTGTGG	7620
3 5	GGATAGTGTG AGGATTGAAT AAGATGAATA CACATGGCTG AGCACACAGC AAGCACTAAA	7680
	TAAGTGCCAG TTTTAATGAT AACGGTGATG ATGATGATGA TGATGATGAT GACGTAACAT	7740
	TGCTTGTGGG ACTCCATACA GCTCAGTAGA TGCTTGCTCA AAGAAGCAAG TTACCAAAAT	7800 [.]
	TTTTGTAATG GTTCTATGAA CGTGAAAAAA GCAGTCAACT TCTCTGAGGA TCAATTTCCT	7860
	TAGTTTCCAA TTAGGAAAAG TCTTCTTAGC TCCAGAGTCC CACAGGGCTA ATGGAATAAG	7920

GAGAGGATAG ATCACACATG TATTATGCAA ACACACTCA GGTGAGCTCT ATTCTTCCTT 7980 CTCAGTTATC CCTTCTGTAG GGACCCCAGT GTCCCTGCT GTCTTTCTGT GTCCTGACCG 8040 GGAAACACAG TGTGCCTTGT CTACTCCATC ACTTGGCCAG CTGCATGCTT TCCTTTGCAG 8100 GCTTGAAGCA AAGCTGGGTC TCGGACATTC TCAGGCACTG ACAAAGCTGT TTAGTTGTTG 8160 5 CTGGGAAACA CTGGGAAATA GCCCTTTTGT TAAACACACA GAAACTAGCC TTCGCCCTGA 8220 GCCAAATTCC TTAAACTCGT CTATGAAATT CCATAACCTG ACTCCTTAAC TGCAGACATA 8280 CCCAGCTAGA ACATCCCTCA TGTCCCTGTC CACCGTGAGA ATGCTGCACT TCACTCTGAA 8340 CCTTTAGTCC TCCTTTTARA TACTGCACAC TGATCACCCT GGTGTTTAGT GCTTTGTTTT 8400 TTGGAATCCC ACCTGGCTCC ATTTTGGGAT GGTTCCGGGC ACTTCCCTAT GGAAATTCCC 8460 10 CTGCTGTCAC TGTCAGAGTG AGTCCAGCAG TGGGTTTAGC TGGATGAAAC ACCACCATGT 8520 CCATTTCCAT TCAGACTAAT GTCAGAATTT GAAAGGCACT ATGGTAGAGT AGAAAGAACA 8580 AGGAACTGTA CTATTTAAAG GGCAGGCAAA GAAAAGGCAT CTATAGCTTA TAAGATGTGT 8640 GGATCTTTGG ATGTGACTTG GCCATCCTGA GCCTAAGTTG TCTTGTAGGA GAAATGGGAA 8700 TGAGAATATT TTCCTCTAGA CATCAAGAGG AAAAGAAATA TAACGTGAAA ACCTTTGTGA 8760 15 ATTGTGAATG TGTTATACAG AGTAGCTAAA AGAATTAAAA AGGGAGTGAC AAAAAAGTAA 8820 AAGGCAGCTG GCTGCTCAGG GCCTCCATGG AGGGAAGTAC CTTGATATGG TCACTGTGGC 8880 TCAGTGACAG CTCTGCAGGG ACAGGAAATT GATTTGTTAG TGCACCCAAA GTTGAATCTG 8940 CTCCTGAGTA CTGATTTATG GGAACCAAAC ACACAAGAGA TGAAGGATGT GTCAACCAGA 9000 ATGTCCAGCA TTAGCTTGTG GGGAAACACA TACTTCCAGT GACTGAAATA CCATCCTGTT 9060 20 ATCAAGAGAT CTGGGAAACT AAAGTACTGA CAAGAGCTGG CTTGATCTGT GGATTTAGAA 9120 CAATGAGAGT TAGGTGGCCT TGAGGGAGAT GATTCACTCT CCTTCACAGA AGAGCTGACC 9180 TCTGGGGTCA ACAGATATAG CACCTCTTTC CCAGGGACGC TACTGAATGA ACAGTGATGT 9240 GTTCTTATAC TCTGGCCCAG ATTTTCTACA TACTTTCTTA GGTTACAACT TTATTTAGTC 9300 ACATTICAGT ACTGGGGATA CTCCTGTTTA TCTTCTTTGG ACTCGAGTTT TTATGGGAAG 9360 25 GTCATGAAAC AGAGAAAAAT ACAATTTGCA GGGAAACTTA CCAAGGCTTG TAAGGTTACA 9420 AGGATTAAAT GAAAACCCTG TGTAAGTCAG TATATAGTGA AGAAGTAAAT TGAGTTAGAC 9480 CARACGCCAA AATGCATCCG CATTAGARAG ACGATARAGG AAGACTCTGG ATTCAGTTCT 9540 GTTCAAAAA CATTTCTGC ACAAATACTA TGTATGAGGA ACTGGGCGTT GGGGAGATGA 9600 TGATGAGTGA GACATGGTTC TTGCTTTCAG AGAGCCTAGA GACCTGGGTG GTAGCAATGG 9660 30 TAGAGATACA TCCAAGACAC AGAAATAGAT ATACAGGAAC ACAGATGATT GAAAGTGATG 9720 CTTGGCAGGG CTTTAAAGAA TGAATCAGAG TTTTTCAGGC AGACGAGGAT CTTCAAGGCA 9780 GAGGGAATCA TATAGATAAG GACATAGAAG AGTGAAATTT CATGAAGTAG TTAAGCATCT 9840 GAAGAAGCAT GGAATTAGTG ACAAGAAATG ATGCGGAAAA GATATCCAGA TCCAATCAAG 9900 AAGGGCCTTG TTGGCATTCT ATGGAGTCTG GACTTTGGCT TCTGGGTCAC AAGTTCTCAG 9960 **+35** ATGGGGTTTT CATATCTATT ATTAGACCTA CTATGTACTG GTCCAGTGGA AGGGAAAGGG 10020 GTTGTCTTAC TGCTAGTGGA GTAGGAATTG GGTATGGACC ACAGCTTGTC TTGTTTCCAA 10080 GTATTCCCTA AGAAATCTGG TCTGCTGATG GGAGATCTAT TTATGGAAAT GTCTTTTCC 10140 CTCAGGAATT TTATGTCGGA AACAGCTGTC ATAGGTGAGG AGGAACTGGT AAAAGTACTT 10200 ANTAGGAGAG TGTCATGGTC AGATTGGTGT TTTGGAAAAG TCAGCCAGGG CAGATTGGAG 10260

5

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AGGTCCATAT	TGGAGGCAGG	aagacttaag	AGACTATTGC	AAAGGTGAAG	ACAAAAGACG	10320
ATAGGGACTT	GCACTTTAAT	TCCAGCCCTT	AGAAGTAGTA	GAAGGTCAGA	AATGAGAATA	10380
TGCATTACAG	AGATAGTTAG	TTGCTATATC	ATTAGGACTT	GGTGATAGAT	TGGATGAGGA	10440
TGCGGTTGGG	TGAGGCAAAG	AGGAGAGTCC	ACATTCCTGG	TCTGGGTAGT	AACAAAGAAT	10500
CTAGCAAGAG	GGCTTGTGGG	GAAAGATGCT	GAGTTACGTA	GCAAGTGCAT	CTGCTTTATC	10560
CTTGTAATGA	ATGGGGCTAA	AGGTGTAAAC	Caaagagtca	TCAGCATTTG	GAGGGTAGAA	10620
TAAATCATCA	GATAACTCAG	GAAGAAGGAG	CAGAAGAATT	ACTGATACTC	CCTGGAAGGA	10680
AAACCGGAAG	TAAATGGGAG	AAACTTGCTC	AAGTGGACAA	AGTTTAACAG	ACATGAAGCA	10740
TGAATTC		•	,		•	10747

- 10 (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6693 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: DNA (genomic)
 - (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

20	GAATTCGGCT	GCTGTGCTGT	CGTACAACAT	GCTGTTTAGG	ATCTTGCACA	TGATAGCTAG	60
	GTATTCTTGC	TTCAAATCGC	AGGCACCCCA	CTTACCAACT	GTGTAGACTT	GATCACGTTA	120
	TTCAACCCCT	GTGTCTCTGC	TTCCTCATTT	TACAAATGGG	GAGAAAAATA	GCATCTATCT	180
	CAAAGTTGTG	AAAATTAAGC	AAGTTAATAC	ATATGTGCTA	CGTAGAACAG	TGCCTGGTAC	240
	ATGGTCAGTT	TTTGATACAT	GTTAGGTATT	ATCATTATTA	TCACCTCCAG	AAACAATTTA	300
25	AACTTCTCAT	ATAAGGCTCT	CCAGACACCT	CTCATTGTCT	TCCCTTCCAA	ATCTGCATTT	360
•	ATCTCTCTCT	CTTTGCAGTC	CAGTGTGAGG	CTTGAATCAC	CTATCAAGCC	TCACCTCCAC	420
	CCCTGTGCTT	TACAAAATGT	CCTAGAGCTT	CTATTTACTC	GTCTCACTGC	TCTGTGGGCT	480
	TTTTCACTCA	AGGGCGTTTG	CATGCTATCC	ATTGCTACCT	GTTTTCTGTT	GCTGGTGTCT	540
	GTCTCCTGCT	CTATCTTTGA	AGAAAAGAAA	CAAGAAAAGG	aataactgag	AAACAGAGAA	600
30	AAAAAATGTC	TCTCCCTTCT	GGTTCTTCCA	GACCACCCAC	TCATCCATCT	TGTTCAATGA	660

ā		CAGCTTCTCI	TCCTTTAATT	AATCACTGTG	GTATATTTAT	AAAGCTTATA	TTTATGAAAG	720
		ACCTTTTAAT	TTTTTAGTTA	TTAAAGCCCT	TTCTCTTTGT	CAGGTTGTAA	CTGAGTGAGC	780
		TCTGGAGTTT	GGAAAGAAGA	TCTTAGAAAT	GGGCCAGAGA	GCTCCTTCTG	AGATCCAAGC	840
r		ACGGAGAATT	GCACCTGCTG	TGCATGGTAA	GAGAGTGTGC	TTGGTAGCTC	ACAAGGGCAA	900
	5	GGTGAGAATA	GAAACTTTCA	TGCCTTTTTG	ATGGGGGTTA	TGAAATCCTA	CCAAGAAACA	960
	:	CCAGGTATCA	GATGTGGGGT	CCTGTTTTCC	CAAAGCCACA	AATGCTTGAA	GGAAGATCTT	1020
		GTGTGATAAA	ATAATTACCA	CATGAACCAA	TCTTGCATGC	ACAGCAATTT	TGAGAGCCCA	1080
		TCCTGGGAGC	TAGGTGTGTA	GTGTTTATCG	TATTGTTGAG	GCTCGTAAAA	ATCTTGTATG	1140
		GCTGCAGGCA	AGCCAAACCC	TTGACAGGCA	CTGCATCTCC	GCTGACTCTA	GAAGACCAAG	1200
	10	CCCAATTTCT	TCCCTGTATA	TAAGGGGAAG	TCTCTATGCT	TGGGGTAGAG	GAGTGTTTAG	1260
		CTCCTTCCCT	TACTCTACCT	TGCTCCTACT	TTTCTCTAAG	TCAACATCGA	ATTTGCCTCC	1320
		TTCATTGACA	AGGTGAGTTT	CTCTCTCATT	GCACTGGTAG	GGCTGCCGCT	GGTCCACTTG	1380
		GGATTGGTGC	AGTCAAAACA	CATGTAGGTT	TGAACCTCAA	GTTTCCATGT	TTACATGATT	1440
		AAAAGGATGT	TTTGTGGAAT	GGTCTCCTAG	GAGATATGTT	AGATGTATGC	TTGTGAATGG	1500
	15	TGTTAATGAC	TCTCTCTTTG	ACAAAGGGTT	CGTGGTCGAC	CTAAAGGTGG	GTCAGTGTGA	1560
		CATTAACATT	TAAGTGCTTT	TTATTCAGCT	CTTGAGCGGA	ATTGGGACTC	ATATCTGTTG	1620
		AATGAAGATA	ATAGAAATGG	GGCTAACTGA	ACTTTCCAGG	GTGCAAGTGA	GAACCCTGGA	1680
		AAGGTCTTCC	TAACCATAGA	AAGGGAGTTG	AGTGTGAACA	TAGTATAGAG	TGTTATTGTA	1740
		GCAGAAAACA	TGTGGTCAGT	CAGTGCCAAA	CATCTTTTGC	TGTCAGAGGG	GAGCTCTGCC	1800
	20	TTCTAATAAT	TTTACATTGG	TACTGGATGA	GGCTAGAGTT	TTTTTATACT	AATATCTCCA	1860
		AAAATCAGCT	CTAAAAAACT	CAGATAAACC	ATTTTTTAA	TTTTTTGCTT	AATCATTAAT	1920
		AGTGCCAATC	CAAGGTTATC	CACAACAAAT	TTCAAATCCA	ATTTTGAATT	TTCCTGATAT	1980
		ACTTTTGAAA	TGTGTGTGTG	TCCTGGGGAT	GCAAACCAGT	TTTTATGGTA	ATATACCTAA	2040
		CAAAATTTTG	GAAGGCAAAT	CTCTTAAATA	CCATGCACCT	ATTTCAAAAC	ATAATTGCAA	2100
	2 5	TAATTCTGTA	TGCGCTTTGC	TATTGGTATT	TGTTTAGTTA	CTCCCTTCCA	AGCCCTCTCT	2160
		GAATTAACAA	GTTGGGTTTT	ATTATGCAGA	TGATATTAAC	TTGATCATCT	TCTTCCTATT	2220
		TCTCTGTCAT	GGTCAGAAGA	TAGGAATTGA	GGTTCTTTTC	CAAATGAGGC	ACAGTTCTCC	2280
		ATGGCTATGA	GACTCCATTT	ATGCATCAGG	AGTAAAGGGG	TCTTGTGTTT	TTAGGTGAGG	2340
		TTCCTGGAGC	AGGATCCCGG	GTACCGCGGC	CGCATCGATT	CGATAAGAGA	TGCCCTCTGT	2400
	30	TTCATTAGCT	CTAGTTCTCC	CCCAGCATCA	CTAACAAATA	TGCTTGGCAA	GACCGAGGTC	2460
		GATTTGTCCC	AGCCTTACCG	GAGAAAAGAG	CTATGGTTAG	TTACACTAGC	TCATCCTATT	2520
		CCCCCAGCTC	TTTCTTTTCT	GCTGTTTCCC	AATGAAGTTT	TCAGATCAGT	GGCAATCTCA	2580
		GTCCCCTGGC	TATGACCCTG	CTTTGTTCTT	TCCCTGAGAA	ACAGTTCAGC	AGTGACCACC	2640
		ACCCACATGA	CATTTCAAAG	CACCTCCTTA	AGCCAGCCAG	agtaggacca -	GTTAGACCCA	2700
₹	3 5	GGGTGTGGAC	AGCTCCTTAG	CATCTTATCT	CTGTGCTGTT	TTGGTTTTGT	ACATAAGGTG	2760
		TAAGCAAGTT	GTTTTTCTTT	TGTGGAGAGG	TCTTAAACTC	CCCATTTCCT	TGTTTTGCTG	2820
,		CAATAAACTG	CATTTGAAAT	TCTCCATGTC	TCGATCGCCC	TTGTTTACGG	Cactgtctaa	2880
		CCTGGATGGG	TGTTTTGTGA	GGTAAAAGAA	GACACTAGAG	CCACATGGCA	TATGGGAAAG	2940
		TCATGCACAC	AAACATGAGA	AAAATGCAGA	GGCCAACCAG	GCAACATTTC .	ACCAGACTGG	3000

	AATCACAGAG AGAGCAAGCA CTTTCCCAGA TGGTGGGGAT GTCATGGAGA AATGGAGAGA 3060)
	CCGGGTGACA GGTTTTGTTC ATTTGAGAAG GCTTTCTTGA AAAGGGCAGT GAGCAAGCAG 3120	,
	GTTGGGAGGA AGAGGTGTGG CATTGAGAAG AAGGGAAAGT ATTGCATGAA AAAGTAATTC 3180)
	TTCACGTGGA ACAGCCAGTA AGGAGGGGCA TGAGTAATAT AGGGTCAGCA GTTACTGGAG 3240)
5	CCAGAATACA GACTTTGGCC TGGGGAGTTC AAGAACTAAG AGTGGTAATA GAGAGTTGGA 3300)
	TATTCCATTT CCCTTCTCTT TTTGTGCCAC CACCCAAAGC TCTGCATAAT CTAAGAAGTT 3360	į
	CCCTTGTTGA CACATAGCTC ATACTTGTGA AGTTGTACAA CAGGATAGCA TAGTGGCCAG 3420	,
	AAGCATGGAC AGTTGAACTC AGATATGCTT GGGTTTGAAT CTTACCATCA CCATTTACTA 3480	1
	GTTCTGTAAT ACAGTGCAAG TTACAGACAT CTCTGCACCT CAGTTTTAGT ATGTCTAAAT 3540	,
10	TGGGGATGAT AATGCCTTCC TTGTGGGGAT AGTGTGAGGA TTGAATAAGA TGAATACACA 3600	1
	TGGCTGAGCA CACAGCAAGC ACTAAATAAG TGCCAGTTTT AATGATAACG GTGATGATGA 3660	i
	TGATGATGAT GATGATGACG TAACATTGCT TGTGGGACTC CATACAGCTC AGTAGATGCT 3720	i
	TGCTCAAAGA AGCAAGTTAC CAAAATTTTT GTAATGGTTC TATGAACGTG AAAAAAGCAG 3780	í
	TCAACTTCTC TGAGGATCAA TTTCCTTAGT TTCCAATTAG GAAAAGTCTT CTTAGCTCCA 3840	•
15	GAGTCCCACA GGGCTAATGG ÁATAAGGAGA GGATAGATCA CACATGTATT ATGCAAACAC 3900	
	AACTCAGGTG AGCTCTATTC TTCCTTCTCA GTTATCCCTT CTGTAGGGAC CCCAGTGTCC 3960	
	CCTGCTGTCT TTCTGTGTCC TGACCGGGAA ACACAGTGTG CCTTGTCTAC TCCATCACTT 4020	
	GGCCAGCTGC ATGCTTTCCT TTGCAGGCTT GAAGCAAAGC TGGGTCTCGG ACATTCTCAG 4080	
	GCACTGACAA AGCTGTTTAG TTGTTGCTGG GAAACACTGG GAAATAGCCC TTTTGTTAAA 4140	
20	CACACAGAAA CTAGCCTTCG CCCTGAGCCA AATTCCTTAA ACTCGTCTAT GAAATTCCAT 4200	
	AACCTGACTC CTTAACTGCA GACATACCCA GCTAGAACAT CCCTCATGTC CCTGTCCACC 4260	
	GTGAGAATGC TGCACTTCAC TCTGAACCTT TAGTCCTCCT TTTAAATACT GCACACTGAT 4320	
	CACCCTGGTG TTTAGTGCTT TGTTTTTTGG AATCCCACCT GGCTCCATTT TGGGATGGTT 4380	
	CCGGGCACTT CCCTATGGAA ATTCCCCTGC TGTCACTGTC AGAGTGAGTC CAGCAGTGGG 4440	
25	TTTAGCTGGA TGAAACACCA CCATGTCCAT TTCCATTCAG ACTAATGTCA GAATTTGAAA 4500	
	GGCACTATGG TAGAGTAGAA AGAACAAGGA ACTGTACTAT TTAAAGGGCA GGCAAAGAAA 4560	
	AGGCATCTAT AGCTTATAAG ATGTGTGGAT CTTTGGATGT GACTTGGCCA TCCTGAGCCT 4620	
	AAGTTGTCTT GTAGGAGAAA TGGGAATGAG AATATTTTCC TCTAGACATC AAGAGGAAAA 4680	
•	GAAATATAAC GTGAAAACCT TTGTGAATTG TGAATGTGTT ATACAGAGTA GCTAAAAGAA 4740	
30	TTAAAAAGGG AGTGACAAAA AAGTAAAAGG CAGCTGGCTG CTCAGGGCCCT CCATGGAGGG 4800	
	AAGTACCTTG ATATGGTCAC TGTGGCTCAG TGACAGCTCT GCAGGGACAG GAAATTGATT 4860	
	TGTTAGTGCA CCCAAAGTTG AATCTGCTCC TGAGTACTGA TTTATGGGAA CCAAACACAC 4920	
•	AAGAGATGAA GGATGTGTCA ACCAGAATGT CCAGCATTAG CTTGTGGGGA AACACATACT 4980	
	TCCAGTGACT GAAATACCAT CCTGTTATCA AGAGATCTGG GAAACTAAAG TACTGACAAG 5040	
35	AGCTGGCTTG ATCTGTGGAT TTAGAACAAT GAGAGTTAGG TGGCCTTGAG GGAGATGATT 5100	
	CACTCTCCTT CACAGAAGAG CTGACCTCTG GGGTCAACAG ATATAGCACC TCTTTCCCAG 5160	
	GGACGCTACT GAATGAACAG TGATGTGTTC TTATACTCTG GCCCAGATTT TCTACATACT 5220	
	TTCTTAGGTT ACAACTTTAT TTAGTCACAT TTCAGTACTG GGGATACTCC TGTTTATCTT 5280	
	CTTTGGACTC GAGTTTTTAT GGGAAGGTCA TGAAACAGAG AAAAATACAA TTTGCAGGGA 5340	

-41-

	AACTTACCAA	GGCTTGTAAG	GTTACAAGGA	TTAAATGAAA	ACCCTGTGTA	AGTCAGTATA	5400
	TAGTGAAGAA	GTAAATTGAG	TTAGACCAAA	CGCCAAAATG	CATCCGCATT	AGAAAGACGA	5460
	TAAAGGAAGA	CTCTGGATTC	AGTTCTGTTC	AAAAAACATT	TTCTGCACAA	ATACTATGTA	5520
	TGAGGAACTG	GCCTTGGGG	AGATGATGAT	GAGTGAGACA	TGGTTCTTGC	TTTCAGAGAG	5580
5	CCTAGAGACC	TGGGTGGTAG	CAATGGTAGA	GATACATCCA	AGACACAGAA	ATAGATATAC	5640
	AGGAACACAG	ATGATTGAAA	GTGATGCTTG	GCAGGGCTTT	aaagaatgaa	TCAGAGTTTT	5700
•	TCAGGCAGAC	GAGGATCTTC	AAGGCAGAGG	GAATCATATA	GATAAGGACA	TAGAAGAGTG	5760
	AAATTTCATG	AAGTAGTTAA	GCATCTGAAG	AAGCATGGAA	TTAGTGACAA	GAAATGATGC	5820
	GGAAAAGATA	TCCAGATCCA	ATCAAGAAGG	GCCTTGTTGG	CATTCTATGG	AGTCTGGACT	5880
10	TTGGCTTCTG	GGTCACAAGT	TCTCAGATGG	GGTTTTCATA	TCTATTATTA	GACCTACTAT	5940
	GTACTGGTCC	AGTGGAAGGG	AAAGGGGTTG	TCTTACTGCT	AGTGGAGTAG	GAATTGGGTA	6000
	TGGACCACAG	CTTGTCTTGT	TTCCAAGTAT	TCCCTAAGAA	ATCTGGTCTG	CTGATGGGAG	6060
	ATCTATTTAT	GGAAATGTCT	TTTTCCCTCA	GGAATTTȚAT	GTCGGAAACA	GCTGTCATAG	6120
	GTGAGGAGGA	ACTGGTAAAA	GTACTTAATA	GGAGAGTGTC	ATGGTCAGAT	TGGTGTTTTG	6180
15	GAAAAGTCAG	CCAGGGCAGA	TTGGAGAGGT	CCATATTGGA	GGCAGGAAGA	CTTAAGAGAC	6240
	TATTGCAAAG	GTGAAGACAA	AAGACGATAG	GGACTTGCAC	TTTAATTCCA	GCCCTTAGAA	6300
	GTAGTAGAAG	GTCAGAAATG	AGAATATGCA	TTACAGAGAT	AGTTAGTTGC	TATATCATTA	6360
	GGACTTGGTG	ATAGATTGGA	TGAGGATGCG	GTTGGGTGAG	GCAAAGAGGA	GAGTCCACAT	6420
	TCCTGGTCTG	GGTAGTAACA	AAGAATCTAG	CAAGAGGGCT	TGTGGGGAAA	GATGCTGAGT	6480
20	TACGTAGCAA	GTGCATCTGC	TTTATCCTTG	TAATGAATGG	GGCTAAAGGT	GTAAACCAAA	6540
	GAGTCATCAG	CATTTGGAGG	GTAGAATAAA	TCATCAGATA	ACTCAGGAAG	AAGGAGCAGA	6600
	AGAATTACTG	ATACTCCCTG	GAAGGAAAAC	CGGAAGTAAA	TGGGAGAAAC	TTGCTCAAGT	6660
	GGACAAAGTT	TAACAGACAT	GAAGCATGAA	TTC			6693
	•		:				

(2) INFORMATION FOR SEQ ID NO:3:

25 (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24979 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear
- 30 (ii) MOLECULE TYPE: DNA (genomic)
 - (iii) HYPOTHETICAL: NO
 - (iv) ANTI-SENSE: NO

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

	GAATTCCAAG	CTTTCTGCTG	TAAGGAGGGA	CCTCAGGGAG	CCAAGGTCAG	CCTGCAGCCT	60
	TTCTGTGCTC	CTTTGCCTCG	CCTGACAGGT	ATGAGGATGA	AATCAACAAG	AGGACTGGCA	120
	GCGAGAATGA	CTTTGTCGTC	CTGAAGAAGG	TGAGGGAAAG	GGGAGTCCTG	AGGGTGGCTG	180
5	TGGACCCAGG	AGGCTGAGGG	GAGTGAGGAA	TCCCTATGGA	TGCTCTGTGA	CAATGGCAGG	240
	GTGGCCTCTA	CGGCCGGCTT	GCTGTGTATG	ATGCCTGAAT	GCGGGGCCCT	TACATTGGAA	300
	CTGACACTGA	Taatgačtct	TCAGGAAGCC	TIGAGTICGT	ATCTCTCTGG	GGTCTGAAAG	360
	TGAAATGAAG	TGAAATGACA	GCTTTTGAGT	GTCAGTTACC	TGTAGCCTTG	GGACCTAAGG	420
	AAGGACCTGG	GGTGTTGGTT	GTGACTGACT	GGGATGTGGA	GGTTGGTGTC	ACATCTCCTT	480
10	CTGGCCAGGA	AAGCCAGGAC	TTGTGGGTCC	TTATTCGAGT	GCGGTGATGA	ATTTTTTAAG	540
	TAAGGAAATA	AACCTAGAGT	GGCTCTGGTC	CTGAGCCAGC	CAGTGAGCTG	TGGCAGGCAA	600
	TGCCTGGGCA	ATAAAGTCAA	ACTGTTCTGC	CTGCTATTCA	GGATGTGGAT	GCTGCTTATG	660
	TGAGCAAAGT	GGACCTGGAG	TCCAGGGTGG	ACACTCTGAC	TGGGGAGGTC	AATTTCTTGA	720
	AATATTTATT	TTTGACGGTG	AGTTAAGCCT	TTATAAGAAC	CTCCTTTCTT	TTCTCACATC	780
15	TCACAAGGAG	TATEGECTET	AAGAGGGGAG	GCCTGAAACC	CAACACTACC	CACTAGGGAC	840
	TCATCTCCCC	AGGTACCCCA	ACTCTGTGGG	CCTGGAGTCA	GCCATCCTCT	CCACCCCAAT	900
	CCTCAGAATC	CCCAGGTCGG	GGTAATGAAG	ATGGAAGGCT	GGGAGAATCC	TGAGTTAGGT	960
	GGAGGCGAAT	GTGTCCCTGG	TTCATGGCTT	CCAATCTGTC	TGGGAAATCA	CCCAGACATA	1020
	TAAGGGGCAA	AACCAACCAG	AAATCTTCAT	TAATTCTGGG	GAGTTGATGG	AGCTGTTAGG	1080
20	AACTCTGTGG	GAGGTGACAG	TGTGAGTCTC	aaggagtgga	CTGACCTTAG	TGATGGGGGA	1140
	TCAAACACTC	CACCACCCGG	CCCTCTTTTG	CCTGTGTCTA	ACTTGGGGGT	ACGTGCTCTG	1200
	GGCCAGATGC	TGTGTTAGAA	GTTTATGTTA	TGGGTATCTC	CATTCTACAG	ATGGGAAAAC	1260
	TGAGGCACTG	AGGGGTTAAA	TTACTTGCTT	CATTACCTAG	CTAGTCAATG	GTACAGCCAA	1320
	Gactcaaaag	TGAGTCCAAG	TGACTCCTTA	ACTAGAGTCC	ATCTACTGCC	TCGGAGTACT	1380
25	CATGTGGTTT	CAAGGAAGAG	GCATGCCTGC	CAAGGAGCCC	AGCTCACTAT	GGTGGCCAAG	1440
	TCAGAGCAAG	GCAGAGTGGC	AGCTGCAGGA	GAAGTGTGAT	GGGGAGATGG	TATCTGAACG	1500
	CTCCAGGTTT	AGGCTCCTTC	CTTCTCCCCT	GGAAGGCAGT	TAAGACTCTC	CCTATTATCT	1560
	CTCATTGCAC	ACAACAATTC	CAAGAGCTTT	TCCCAAGACT	ACCTGGCCCA	GGCTTCTGGC	1620
	TTCCCCCGAG	AGCCTTGAGG	GAGCAGCAGA	GGAAAACTGA	GGCCCCCAGA	ggagaatgga	1680
3 0	AGGAGTCAGC	CTGTGCGCCA	TGCCTCGCAG	GAGCTGTCTC	AGGTGCAGAC	TCACATCAGC	1740
	GACACCAACG	TCATCCTGTT	CATGGACAAT	AACCGTTCCC	TGGACCTGGA	CAGCATCATC	1800
	GATCGAGTGC	GGACCCAGTA	TGAACTGATT	GCACAGAGGA	GCAAGGACGA	GGCCGAAAGC	1860
	CTGTACCAGA	CCAAGGTGGG	CGTGGCCCAG	ATCTGGTGCC	CAGAAAAACA	GATTCTTCCC	1920
	AGAATTGTCT	TTCTCTTATT	GCATTGTCTT	TCTCTTATTT	CTGAAGTAAA	ATGTGTTTGT	1980
35	TATACAAATT	CTAGAAATTA	CATGTAAAGA	TTACCCATCT	CTCACTACCG	CTATTAATAT	2040
	GTTAATATCT	CTTCTACCAG	TȚCTTTGTCG	CTATTAGGCT	agtggaaaag	TGATTGCGGT	2100
	TCTCGCCATT	AAAAGTAATG	ACGAGAACTG	Caattacttt	TGCAAAAACC	CAATAATGTT	2160
	TATTGAGAAC	TCATATGTGT	TAGGCACCTA	GCAAAGTGCT	TTACTTATTT	ATTATTATTT	2220

•	CATTCAGTCC TTACAACAAC CAATGAGGTA AGAATTTCGT TATCACCATT TTATAATAGA 2	280
	TAGTAGTGTG TGACATTACT TAATTTCCCT AATGCCTTGT AGCTAGTAAA TGCAGAGCCA 2	340
	GAGCTTAATT AAAATTGGTT TGTGTCTACA AACCCATTCC CCTCACCACT AGAATGATTT 2	400
£	TTATTCTTTT TTCATAATGG TATCTATTAA AATATATTTT TTTTACTTTT TTTTTTTT	460
5	GAGATGGAAA CTCACTCTAT CTCCCAGGCT GGAGTGCAGT GGCCTGATAT CAGTTCACTA 2	520
	CANACTCCGC CTCCTGGGTC CANGTGATTC TCCTGCCTCA GCCTCCTGAG TAGCTGGGAT 2	580
	TACAGGCACG TGCACTACAC CAAGCTAATT TTTGTATTTT TAGTAGTGGC AGGGTTTTGC 2	640
	CATGTTGGCC AGGCTATACT TTCTCTTTTA CTTAACACAT ATGGATACTT TTCTGTGACA 2	700
	CTAAATAACC TGCCGCATTT TTAACAGCCT GGTATTATAT TGTTAGACTA CACCCTCCCT 2	760
10	TATTAGATCA ACTOCTTTGT GGCTAAGTTG TTGGGCACAT CTTTGGTTAC TTCTTTTCCA 2	820
	TAAACTGACC TGGATCTTTT TGGATGGTGA AGCCTCTGGT TGAAAGGGTG TGGCTGACAG 2	880
	TCCAGTCACT AAATTCTGAA CAACTAGCAT TGATGACTGG CTTTGAGGAT GATCTGTGGC 2	940
	CAACTCCAAT CCTGGCTGAC CTCTGTCCCA CGGTCCTGCA CAGTGTTCTG GGGTGGAATG 3	000
	GATTTCGACA TTAGACTAGG AAGCCAGATG GCCAACAGTG AAAAATAGCA GAGTGTACCA 3	060
15	GATTCCCTTG CAAGTCGATG CTTCTCCTAC CCACTTCAGA GCCCTGTGCC TGGGGGGTGG 3:	120
	AGTTCTGACT AATGGGGCAA TACAGAGACA GAAACAGAGA TGGAGGGGAA ATGAGACTGA 3	180
	ACCTGGAGCC AATGGAGGC CTCTGAGGAC ATGAGGTCTG CTTGACTGCT AGGGAGATCA 3	240
	TCCTGGAAAA GGGTGGGAAG CTATATGGTG GGTGGAAAGA GTGAGGGGGT CTCAGTGTGG 3	300
	GTAAGGACCA ACGTGAAGGC TTAGATGTGT GAAAAGGTGG TAGAAGGGCA TCACAAAGCA 3	360
20	GGTTTGTCTG GCCTGGGATG AGAGTCTGCC CAGAGACTGG TGGGAATGCG GGAGGCTTGG 34	420
	GATAGTGTGA GTGTGTGCAT GTATACATGT GTTTGCAGCC TGGGTGAGGG AGGTTTGGTA 34	480
	TAGCTGTGAG TATGCATGTA GGGGTGACCA CAGTGCAAGG TGGGTGGGAA TCTCCCAGGG 3	540
	GAGAGCAGCC CAGACCTACT CCTCCTGGAG GGGCTTGTGG TGGGCAGCAC ATGCTGACTA 3	600
	TGATGCTCGC TTTGGCCCCA GTACCAGGAG CTCCAGATCA CGGCAGGGAG ACATGGAGAT 36	660
25	GACCTGAAGA ACAGCAAGAT GGAGATTGCA GAGCTCAACC GCACCGTCCA GAGGCTGCAG 3	720
	GCAGAGATCA GCAACGTGAA GAAGCTGGTG GGACGGGTGC TTAGGGAGGG CTGACCAAAG 3	780
	CCCTGCACCT CCTACAATGC CCTGCCAGAT CGAGCTCTGG AAACTTAACC ATTAAATGGT 38	340
	CTCCAACTGT CTCTGGAGCA GATTGAACAG ATGCAGTCAC TCATTTCGGA TGCTGAGGAG 39	300
	AGAGGCGAGC AGGCCCTCCA GGATGCGTGG CAGAAGCTGC AGGACCTGGA GGAGGCCCTG 39	960
30	CAGCAGTCCA AGGAGGAGCT GGCCCGGCTG CTGCGTGACT ACCAGGCCAT GCTGGGGGTC 40)20
	AAGCTGTCCC TGGATGTGGA GATCGCCACC TACCGCCAGC TGCTGGAGGG CGAGGAGAGC 40	080
	AGGTGGGTCT GGCAGCTGTG TTTCTGGGGC TAAGGCTTGA GATGCACCAT GAAGCTGTGG 41	140
	GACTGGCTAT TTGGAGAAAA GATAAGCCCA CCTTTTTGGG AAGATTGGTA GCCAGGTGAG 42	200
	CAGAAACATT CCAATTAGAG GCAGAGGCTG TGTGAATGGA CAAGCCTCTT CACACAGGGA 42	260
* 35	GAAGTCATTG TTATCATTCC TCCACCTCCA AGTAGAATGT CCTTATACCC CAATCCAAGC 43	320
	CTCTGCAGCT GGTATTCACC CCCAATGCTA AAAGGCTTCA TGAAAACCCT GAAATTTCTC 43	380
á	TCTGCCCCAC TGGCTTCCTG ACCTCTGCTC ATGCACACAC ATTTCCCTAA GGCTTGGGGA 44	140
	CACCTCTGAT CCAGATGTCT GTGGCCACAG CCTTCTCTCC TCAGGCCCTG TGGGTCTGGC 45	500
	TGACCGTGTG CTTTGGTTTT ACAGGATGTC AGGAGAGCTG CAGAGCCATG TGAGCATCTG 45	560

	TAAGTAGCAG	ACCCAGGGGC	AGAGAGAGGC	TGGTGGTGCT	GGGTGGAGGG	AGGGCCAGGA	4620
•	GGTGGCCAGC	AGAGAACGGA	AAGTCTGGCA	TTTTAGCTTC	CAGTCCTGTG	CAATAGACAC	4680
	CAAAGTAAGC	AAGTGTAATG	CAAAGCCTGG	AAGAATTCAT	TTCAAATAAA	TGGTTATGAT	4740
	TTCAGGTCTG	CTTATCTTAA	TTGTTATGAT	GCCTTTTTAT	TARATGATGC	CTAGGAGGAA	4800
5	TCAGCAGCGG	CTAGAACTCT	TTAGGGTACA	TATTCAATAA	ACAATGTAAG	TGTGTTGCTG	4860
	AGAGGAACCC	TGGCATCCCT	TTGTAGTATG	AAGAATACTT	TTCAAGTAGG	AACACTTTCA	4920
	ATTTTCAATG	TATCGGGTTT	GCAAGTCGAT	GCCACGGGTG	ATCGAGGATG	GAGGAGGCTG	4980
	CAGGTGCAGG	GCGGGTGCAG	CTGGGGGCCT	TGCCTGTCCC	TCTGACCCCG	TGTGCACTGT	5040
	CCCTACCCAC	AGCCGTGCAG	AACAGCCAGG	TGAGCCGTCA	ACGGCGGCGC	GGGAGGCGGC	5100
10	GGCAGCTACG	GCTCAGGAGG	CTACGGCGGC	GGCAGCGGTG	GGGGCTATGG	CGGCGGAAGA	5160
	AGCTACCGCG	GAGGCGGGC	ACGAGGCGGG	AGTGGAGGCG	GTTATGGCAG	CGGCTGCGGC	5220 ⁻
•	GGCGGTGGCG	GGAGCTACGG	AGGGAGCGGC	AGAAGCGGCC	GCGGATCCTC	GCGCGTGCAG	5280
	ATCATCCAGA	CCTCCACCAA	CACCTCCCAC	AGGCGGATCT	TGGAGTAGAG	GCCTCGTTTC	5340
	TGCCACACAT	CACGCCTGCC	CCTCACCGAC	CTCTCCTCAA	ACTCCTCCCC	TCCACGCCCT	5400
15	TCCTAATCCC	CTCTCATTCA	CTTTTCTTAA	TGGGTCTCAG	CAATTTTGCC	AATAAATTCG	5460
	ACTCTAATGG	GGGAAGCAGG	GTGGATAAGT	CCAAACAGCA	GATCTCTCTT	TTGGAGGGCA	5520
	CTGGCTTGCA	GTCAGATTCA	CAGCTAGGCA	CATTCTCACT	CAGACCCCGC	TCTGCTGGCC	5580
	CTCCTCCTCT	TCCTGCTCCC	ACCTTTTTGG	AAGATCGGTA	GCCCAGGGTG	AGCACAAACA	5640
	TTCCAGTTAG	AGGCAGAGGC	TGCGTGAGTT	GGCAAGGTAG	GGAGAAGTCA	TTGTTATCAT	5700
20	TCCTCTGCCT	CCAAGTAGAA	TGCCCTTATG	CCCCAGTTCA	AGCCACTGCA	GCTAAGTATT	5760
	AACCCCCAGT	GCTAAAAGAT	ACCAGGCATC	TAGTTTAGCA	ATGGAGGGAA	ACAGAAACAG	5820
	CCTGTAGAGA	GCATCGACAA	GGCGCATAAT	GGAGAGTTGG	TATCTCAACC	CCAAGCTCCC	5880
	TTTGCTGTAC	CIGGGCCIGC	TCTGTGAACA	AGAATCCACG	CCCCCTGCC	CTGCTGGGAC	5940
	CACATAATGA	TCCCTTTGGG	GAAGTTGCTG	ATTGCAGGGC	AAGCTAGCTT	GGTAAGGAAA	6000
25	ACCTCTGCAC	CAGCGGCCTA	TICCIGCCIC	TTGGTCCATA	GCCTCATACA	CTCATGTTGA	6060
	TGGATAGTAT	AGATTTGCTG	CCCACACCAG	ATATCTGTAA	GGCATCACTG	TCCTGATTCT	6120
	GAACCTCTGT	TTCAGGAAGC	ATTCTCCCCT	GTGTAAACAA	CTCAAGGTGG	AAGTATTTCA	6180
	GAGGGCATAG	GGTCATGAAT	CCTTACCCAA	AGGAAGCCTG	TTTTAGCAGT	GGATGCAGGA	6240
	GTTGATGAAC	agacaagcaa	GTTCTGCTTC	TGTCCTGTTT	CCTCCTGACA	GCTCCATTCT	6300
30	TTTGAAGCCT	GACCCTTCCT	AAGCTCTGCA	TCATAACCAC	TCTGAGAATT	GCCCCATTGG	6360
•	TGGGCATGTG	AAGCCAGCTC	TGTTCCATCC	AGGTGCCTCG	GGCCTGAGGA	GTCTGAGGAT	6420
		TCTTGGAAGG					6480
	CAAGGCATTT	TAATGCCTTT	CCCAGACCTC	TCTAGTCCCT	CCTGCCTTCT	GTTCTCTCGA	6540
	CAGCTGTGAG	CCCTTTAGAG	AGAATATGAC	TCTTAATTTT	GAATCTTATG	TAAGAGGCTT	6600
35	GAGATGTGCT	GGAGAGGGCA	GGAAGAGGAA	AGTATCAGGC	CTTGAGAGAG	GGAATGTAGC	6660
	TTTGCTTCTA	TGATCTGGAG	TCACCTTCAC	TTGCTAGCTG	AGTCCTAACA	CAACTTCCAA	6720
	•		•			CTGTTCTTGT	
						TGGTATTTCT	6840
	ATAATAAATT	TTCTGTGATC	TCAAATTGCT	GTTTGGTCAG	GAGATGCATT	ATTTCTTCTT	6900

CTTCTTCTCC TTCTTCTCT TTGCCTTCCT CCTCTTCTCC CTCCTCTGCT TCTCCTTGCT 6960 CCTCTGGCTC CTCCTCCGCG CTCCTCCTCC TCCTCCTCCG CGCTCCTCCT CCGCGCGCTC 7020 CTCCTCCTTC TTCTTCTTCT CCTTCTCCTT CTCATCTTC CATCTTCATC TTCATCTTCT 7080 CCTCTTTTC TTCTTCCTC TAAATAAAGA TGGGGTCTCA CCATGTTTCC AAGGCTGGTC 7140 5 TTGAACTCCT GGGCTCAGGT GATCCTCCCA CCTTGGCTTC CCACATTGCT GGGATTACAG 7200 GTGTGAGGTG TGGTACCTGG GCTATTTCTT TAAAAATTTC TGCAGACCTC TGAAATTATT 7260 TATATTTGGG AAGTTAAAAT TTCTTCTTAT TTTTTATTGT ACAAGTAATA CACAGTCTTG 7320 AAGAATCTTA CAGACATAAT CTTATTAATC CTTAAAGTGG CTGATCATCC AAAAGTCAAT 7380 TATACATTTG TTCAATGAGC ACTTATTAAG CTCCTACTGT GTGGCGGGCA GTGGCTTAGG 7440 CACTGGGAAT GCAATGTTGA ATGAACATGT TTCTGACTCT TAAGTTGCTC ACAACTAAAT 7500 10 GACCTCTAGC CTGAGAGTGG AAGCAAGGCC TATCCTGAGG ACACAGGCAG ACCCCCCAAA 7620 ACAGGAACAG GTGGGACTTA CGACAGGTGC CAGTGCTGGG GAAGGGACGT TTGGTTCCAA 7680 CAGACTCCTG GAGGACTGGG ATATGGAACA GGGCCAAGGA AGAGAGGTGT GGGTGGGGAG 7740 15 ATGAGGGAAG GGCCCTCCAA ACAGGGGGAT AGTCTGCTCA GAGACTCAAA ATAAGAGAGA 7800 GTGTGGGGT GAGAAGGAC AGCTGGACAG GAGAAACTGA GCTAGGGAAG GAAGGGGCTG 7860 AGGCCACAAA CTGAGTGGGG TCATGGGCAG AGACATCTTC AATTGATGCC TTGAGGGAAG 7920 CAGAGATGCA GAAATTCCAT AATGGAGCAA GTTAAGCCAT CACCTCATCC TATGTGGTAG 7980 TTCTCAGTCC ATGTAAAAGA ATCACATAAA AGATGTGATC TACTTTCTAA TTCCCTGGAG 8040 20 GACTTTGCAT GCAAATTTGG ATATGGGATT CATTCGAATA TGACAGGAAC CCCATATTGA 8100 TANGACACTG TTGCTCCCGG GTGGGCATTG TTCAACTCAA GACTTGATGA CCCAGATAGG 8160 TGTGTCTTTG CAGTTAGCTG TCACATGTCC CACCGTTGAA AGGTGGGCTT CTCCTCCACA 8220 TGTGCAGGGC TCTCTGCCTG CCTTTCCCTT TTCTCGTGTC CTCTGACAGC CTGCTGCCAG 8280 GATAGATGAG ATGGGGAGAA ACTTCTCAGA GAGAATAGAG GGGTGTGCAT GGAAACAGAG 8340 25 TGTCTTATCA CTATGGGTTG ATATGATGTT TGCAGTTAGC TGCCACATCC TCCCCAAAGA 8400 CTTCTGGAGG GCATGCCTGG GAACACAATG TTTTATTCAT ATGGGTTGCT GTCCTATTCC 8460 ANTGANTCCC ATATCCAAAT TCCATCAATA TCGCCTTCAG GAAGCTACAA CATATTCGGC 8520 TCANTATAAG AAGCACCTTT CTATGATCCT GACATGGGAG AGGCTACCCT GGGGAGTGAT 8580 CAAGTTTCAA GTCAGAGATT GGCTAACCGT TTGGCAGGAA CGTTGAGGGC GGGAGTGGAG 8640 30 ATGGGTGGGG ATATGGTATG GAGGCATCTC ACTACTTTGC TGTACTAAGA GTTCACATGG 8700 CGAAACCTGA GAAAAAAAT TCTACTCTCT GTGTTATATG GGAAGAATAA GGTCAGGTGC 8760 CAGTGAAAGC TAAAGTCACA AAGAAGCCAA AGGCCCTAGC CAGAACTGTT AAATGAGGCT 8820 AAGTTTTCTG GCAGCACAGG GTCTATTACA GGGTGTGAGT TTGATTATCC CTGGGATCAT 8880 GCATGTGTGA TACTCTAATG GGATCCACGT TGGCTCTGAG AAAACACGCA AGGATAAGGC 8940 - 35 CAACCACAGC TCTCCTTTCC CATCCTCTCT TGGGAACAAG TTGAGATTGT CCCAGAAAAT 9000 GTGGCCCTGA CTTATCTCTT CCGAATTCCT TGATTTTGTC CTGTCATGGA GGCCTGGGGG 9060 ACAGATGGAG GGAATCATGT GCCTGAATCT GAAGAATATT GGAATAGAGA TTCCACAAGG 9120 TAGGGGCAGG AGAAATAAAG GACAGAAAGG AGAGGAGTTG GTCAAAGAAG GCATCTCAAC 9180 GTCTAAATGA GAAGTCTTAA TTCGATGTTC AGGGAAAGAA AGAGTAACTT TAGGGACCTA 9240

	AACAAGGAGG	ACTAGCACTA	AGACACTGAA	GAGATITCCI	GARATAGACA	MINITIOONI	,,,,,,
	CAGAGACAAT	GAGAAATCCC	ATCAGGAGAA	AATGTCTCTC	ACTITCAGCT	CACCCCAGTG	9360
	AAAACAACAA	GCATTCTATA	AACCATGTAG	GAAATGCCCA	CACATGCATT	ATCTCACCTG	9420
	AGTCCCACTC	ACCTGGGAGT	GCGGAGACCA	GGTGTGGGGG	TCTGCAGTCC	TTCTAGGGAC	9480
5	CATGGAGTGC	TCCATCCCTG	CCCCTAATCA	ATGCTATTCC	CACAAGGCAG	ATACTCAGAG	9540
	GGAGAGCCAA	GCAGGCTCAT	TGCAGTGCAA	TAAAGCCAAG	AGGCTGGCAG	GAGGGAGCAA	9600
	ACACCCGGGT	TGGTGAGAGT	CCCAGGGAAA	GTCTGCCAGT	CTGCTCTTTG	CTCTGAGAGG	9660
	CAGGGTGGCA	GGGTTGGGGC	ACTCTGGAAA	TATAAATTTA	GTTCCACCAG	CTTCTCATCC	9720
	ACAGAGATTT	TGATCTGAGG	ACATGGTTAA	CTGGAGGAGC	AATCATTGAC	TCAGTAAAAT	9780
10	TCTAACTGCA	TCTGACCTTA	GACAAGGTGT	GCGTTTCTGG	GCTGGGAAAG	TTCCTGGTCT	9840
	GAGGAAGAGT	CTCTTGAGAA	TGTCATCTCT	TTTCAATTAC	CCAGCCTTTT	GGCCCAGAAT	9900
	GCATCTTCAA	ATTAATGAGC	CATTTGCTGG	TTAATTTGĠT	CCCAGGGAAA	AAAGTCCAGC	9960
				•		TGGTTTTATC	
	AGCTTTCCGA	TCAGTCTGGG	AAATGGGATG	TCTTCAAGGC	TGATCAATGC	TCTGTTGAAG	10080
15				_		CTCCGAGTGG	
	CCCAATCCCA	AGCCTGGAAG	GGCTTCCAGG	GGGCTCTAAG	TGTGCATTCT	GACCTCCACA	10200
	CCTGCCCCTG	TGTGCTCAGC	CCTCAGTGTT	TGTGCTCCCC	CTGCAGAGCA	GCTCTGCAGT	10260
						AGCCCGGTGA	
	_	•				AAGTCCGTGG	
20						GCTTAAATGT	
			•	•		GTTTGTTTTT	
	TTTTTTCTGG	TTATTATATC	AGCTTCTGGG	TTCTCTCAAA	TGCAAGAGTG	AGGGAAAATC	10560
	TICCTITITI	CCTTTTTTGA	GATGGAGTCT	TCAGCATCAG	TAGCCCAGGC	TGGAGTTCAG	10620
		-				CTCCTGCCTC	
25	AGCCTCCCGA	GTAGCTGGGA	CTACAGGGCC	TGCCACCACA	CTGGGCTAAT	TTTTTTTGTA	10740
	TTTTTAGTGG	AGACGGGGTT	TCACCTTGTT	AGCCAGGATG	GTCTCGATCT	CCTGGTCTCG	10800
						ACGCCTGGCT	
		•		_		CATAGAGCCT	
		•				GTGCAGAATC	
30						CCACCGCCTC	
						CCAGCCTTGC	
						GCTGCCCCTC	
						CCTGGCTGCT	
	TTCAACATCT	CGCCCCGCCT	CCCCATTGTC	TGTGATTTCT	CTTCACTCCA	CCCAGGCCTC	11280
35		-		• •	· ·	CCCTCAGGGT	
						CAAAAGCTCT	
•						AGCCTCATCT	
						GGTGCTCTGT	
	CATCAGCACC	GCTGGGGTAA	CTCTCAAGTA	TAAGGGGCCA	TGTGGGATGC	TGGGAGGGCA	11580

TCAAAAGACA CAGGGGACTT AGTCTTGCTT TCCAAAGGCT TCCAGAGTGA TTGAGGGGCC 11640 AGGAAACACA CAAGCACATG CATGAAAATG AGCCAACAAA TGCATCAATA TGTACTAAGT 11700 CTGGCAGCAG CCGAGCTTGG AAAAAGAGAC CAATAGAGCA CTTGCCCGAT GTGGACTGAG 11760 CAAAACTCCC TGGAGGAGAT GAGATCTGGA CCTGTCTGCT GCCTGCTTTG AGTGAGAGAT 11820 5 GAAGGCATTT GCCCACAAGC CCTGATGGAC CAAAAACAGA TTCAGGACCA AATGCTCAGC 11880 CATTGAGATC TTTGGTGCCC CAGAGCTTGA CTATGGGTAG GGATTTGTGG CAATGCCGAG 11940 GCAACCAGAA GACCTTTCAG AAAAGAGAAG AGTAGAAGTG GGCTTGGAAG ACAGAGAGGA 12000 ACAGGGATGG AAAGGGAAGA AGAGGGTGAT CAGTCTTGGG CAAAGCACGA GAGCTGAAGG 12060 GGTCAAGGCT GTGAGGCCGG GGAAGTGGGT GAGCAGGGTA AGATGTAGGT GGTGCTGGTG 12120 10 GTGAGAGCAG GCCAATGACA GAAGGAGCCC ATGTGATGCG GCGGGCTTGG ACTCTGGAGT 12180 GAGGCACGTG GCTTGTCAGT TACCTGCTGG ATGACTTTGG GCACATATTT CAACCTCTAT 12240 ANACCTAAGA TGCCTTTTCT TTAAAATGGG GCTAAGAGCT CCCACGACGC AGACTTTGTG 12300 TGGTATTTAA ATGCAATGTG GCTCCTAACA GCATAGTTGC TGCGTGTAGA TGTTAGTGTC 12360 TCTTTCTTTC TCATTTTGTC TTTATTTCAT AAATGCACAG TCACTAAGTA AGAAAGGAGA 12420 15 GAGTGTGTGG CTCACACTTT CCTGCATGTG GTTCTTCATA TCCCACACAC CACACTGATC 12480 CTGGGGACAT CACAGGAGAT GACGGGCCTG GTCTGGCAGC ACTGCAGCTC CAGCTCTGTT 12540 GGGCTGCCTC GAAAGTGGGC AGTGGAAAAA GAAAAGGAGT TTGATTCAAC AATTGGAAGA 12600 TACTTTTTTG GGCCTGTGCT CTCACTTCTC TGTGAGGCAG GTTAGATGAT GTGACCTTTG 12720 20 AGGCCCCATG GATGAGAACA TTCTGTAATT CTCTGTGTAC TTGTTTATAG GGCCCAGTTC 12780 CACTTGCCTG TCTTTGAGCC TCTTCCCGGT TCAGGGAGGA ATGTCACTTG AATTGAAATC 12840 AGAAAACCCA GATTCTGCTT CCAGATGTGT CTTTTCCTAG CCGGAGTGTC TAGAGGAAGC 12900 CACTTAATCT CTGAGAATCA GTTTTCTGTT TCATGAAATG GGTTGAGAAC AGCTTGATTG 12960 CCTAGTTCTC AGGGCTCTTG TGGGATGCTC TTTGCATATG TGTTTGGTGG GGTGAGCTGT 13020 25 GCAAATGTAA GCTATGGTGA GGTTTATGGC ACTTATTCCT GCTAGTCCTG CATTTCTCCC 13080 TTCTCACAGG AGCACCTGGG GTATGTTTTG CAGCTAAGTT GTCTACCAAT TCCCTGACCA 13140 TTCATTCAAA CCTTTGATTT TTCTGTATGT CAGTTTCTTA GTTCAAAGAT GGGAGTGTGG 13200 ATCACTGCCA AGGTCTGTTT TTGGCTGGCA CACACATGCA CACAAACATG TGTGCACACA 13260 AACATGTGTG CCCAAACATA CTCACACCCC TCCAAAATGC TAGAAGGAAT CGATTGTGCA 13320 30 GAACAATATG TCTCATGAGG GAGTATGCTG AACTAAAATA ATTTTGATTG CTTGTCAGAA 13380 ANTGATTAGG CARCAGTCAT TACCATGCCA AGACTGTCCC AGTCTCCATT GTTCCTAACA 13440 AGACCTGAAT TACTCATTCC CTAAAGAGAT GGTTGGTTTA GCAGCCGAAG GATTTTAGTG 13500 CTAGACAGAG TCCCAGACAG CAGTGCCACA GTGATGGCGA GGGAGAGGAG TAGCAGGGGA 13560 GCGGTGAGGG GCACTTTCTG GAGGAGGGTA TAGGGCAAAA ACTGGGAGGA GAAGAGGGAC 13620 - 35 ARGGTTCAAT AGCGGAGTGC AATGGAGAGG ACCGACACAG CCAGCCCGAT TCAGAGCCAC 13680 AGAGTAATGG GACCAGATGA TCTTCACAGA CTCCCTTTCT CCCATAGATC TTGCACACCA 13740 TAGTGGAGAC TTCCCATGTA CATCTATGGT TTGCCACTTA CAGAGTTACT TGGAGCCAGC 13800 TGAAGTTAGA GCTGGCTTCT CCCCTTTGAG TCTTCAATTC TGTGTTTATG TGCAGGCCCG 13860 GGGACCATGC CAGGCTTCTA AGAAGGTCTT CGAATGAAAG TCTGCTTGGG CTCTAGTGTG 13920

TCCAGATCTC AGTGCCACTA TTATCCACTG ATATTGATCA AGTGCTGCTC TCCAGGAAGA 13980 CCCCTGAGGT TTCCTGGTCC ATTGCAATGC ATGCTGGGTA CTCTTGCACT TGGATGGAAG 14040 TAAAATCTCC TCACTAAACT CTGTGCCACC AAAATCTCCT TCTCAGTGTG AATTGAAGAA 14100 ACATTITCCA AGACTIGCAT GIGCCAGGAG CCAAGGACIC AGAGIGATAA AACAGCETIC 14160 TGCCCTCAGA GCTCTCTGTG GTGGGGCGCT TCCTGTGCTG TCTGGCTTTA CACACAGCAG 14220 5 GCAGAATGAC TTGAATTCGG CTGCTGTGCT GTCGTACAAC ATGCTGTTTA GGATCTTGCA 14280 CATGATAGCT AGGTATTCTT GCTTCAAATC GCAGGCACCC CACTTACCAA CTGTGTAGAC 14340 TTGATCACGT TATTCAACCC CTGTGTCTCT GCTTCCTCAT TTTACAAATG GGGAGAAAAA 14400 TAGCATCTAT CTCAAAGTTG TGAAAATTAA GCAAGTTAAT ACATATGTGC TACGTAGAAC 14460 AGTGCCTGGT ACATGGTCAG TTTTTGATAC ATGTTAGGTA TTATCATTAT TATCACCTCC 14520 10 ARATCIGCAT TIATCICICI CICTITGCAG ICCAGIGIGA GGCITGAAIC ACCIATCAAG 14640 CCTCACCTCC ACCCCTGTGC TTTACAAAAT GTCCTAGAGC TTCTATTTAC TCGTCTCACT 14700 GCTCTGTGGG CTTTTCACT CAAGGGCGTT TGCATGCTAT CCATTGCTAC CTGTTTTCTG 14760 TTGCTGGTGT CTGTCTCCTG CTCTATCTTT GAAGAAAAGA AACAAGAAAA GGAATAACTG 14820 15 AGARACAGAG AAAAAAAATG TCTCTCCCTT CTGGTTCTTC CAGACCACCC ACTCATCCAT 14880 CTTGTTCAAT GACAGCTTCT CTTCCTTTAA TIAATCACTG TGGTATATTT ATAAAGCTTA 14940 TATTTATGAA AGACCTTTTA ATTTTTTAGT TATTAAAGCC CTTTCTCTTT GTCAGGTTGT 15000 aactgagtga gctctggagt ttggaaagaa gatcttagaa atgggccaga gagctccttc 15060 TGAGATCCAA GCACGGAGAA TTGCACCTGC TGTGCATGGT AAGAGAGTGT GCTTGGTAGC 15120 20 TCACAAGGGC AAGGTGAGAA TAGAAACTTT CATGCCTTTT TGATGGGGGT TATGAAATCC 15180 TACCAAGAAA CACCAGGTAT CAGATGTGGG GTCCTGTTTT CCCAAAGCCA CAAATGCTTG 15240 AAGGAAGATC TIGIGIGATA AAATAATTAC CACATGAACC AATCTIGCAT GCACAGCAAT 15300 TTTGAGAGCC CATCCTGGGA GCTAGGTGTG TAGTGTTAT CGTATTGTTG AGGCTCGTAA 15360 AAATCTTGTA TGGCTGCAGG CAAGCCAAAC CCTTGACAGG CACTGCATCT CCGCTGACTC 15420 25 TAGAAGACCA AGCCCAATTT CTTCCCTGTA TATAAGGGGA AGTCTCTATG CTTGGGGTAG 15480 AGGAGTGTTT AGCTCCTTCC CTTACTCTAC CTTGCTCCTA CTTTTCTCTA AGTCAACATG 15540 AGTCGACAGT TTAGTTCCAG GTCTGGGTAC CGAAGTGGAG GGGGCTTCAG CTCTGGCTCT 15600 GCTGGGATCA TCAACTACCA GCGCAGGACC ACCAGCAGCT CCACACGCCG CAGTGGAGGA 15660 GGTGGTGGGA GATTTTCAAG CTGTGGTGGT GGTGGTGGTA GCTTTGGTGC TGGTGGTGGA 15720 30 TTTGGAAGTC GGAGTCTTGT TAACCTTGGT GGCAGTAAAA GCATCTCCAT AAGTGTGGCT 15780 AGAGGAGGTG GACGTGGTAG TGGCTTTGGT GGTGGTTATG GTGGTGGTGG CTTTGGTGGT 15840 GGTGGCTTTG GTGGTGGTGG CTTTGGTGGA GGTGGCATTG GGGGTGGTGG CTTTGGTGGT 15900 TTTGGCAGTG GTGGTGGG TTTTGGTGGA GGTGGCTTTG GGGGTGGTGG ATATGGGGGT 15960 GGTTATGGTC CTGTCTGCCC TCCTGGTGGC ATACAAGAAG TCACTATCAA CCAGAGCCTT 16020 35 CTTCAGCCCC TCAATGTGGA GATTGACCCT GAGATCCAAA AGGTGAAGTC TCGAGAAAGG 16080 GAGCAAATCA AGTCACTCAA CAACCAATTT GCCTCCTTCA TTGACAAGGT GAGTTTCTCT 16140 CTCATTGCAC TGGTAGGGCT GCCGCTGGTC CACTTGGGAT TGGTGCAGTC AAAACACATG 16200 TAGGTTTGAA CCTCAAGTTT CCATGTTTAC ATGATTAAAA GGATGTTTTG TGGAATGGTC 16260

TCCTAGGAGA TATGTTAGAT GTATGCTTGT GAATGGTGTT AATGACTCTC TCTTTGACAA 16320 AGGGTTCGTG GTCGACCTAA AGGTGGGTCA GTGTGACATT AACATTTAAG TGCTTTTTAT 16380 TCAGCTCTTG AGCGGAATTG GGACTCATAT CTGTTGAATG AAGATAATAG AAATGGGGCT 16440 AACTGAACTT TCCAGGGTGC AAGTGAGAAC CCTGGAAAGG TCTTCCTAAC CATAGAAAGG 16500 GAGTTGAGTG TGAACATAGT ATAGAGTGTT ATTGTAGCAG AAAACATGTG GTCAGTCAGT 16560 5 GCCAAACATC TTTTGCTGTC AGAGGGGAGC TCTGCCTTCT AATAATTTTA CATTGGTACT 16620 GGATGAGGCT AGAGTTTTTT TATACTAATA TCTCCAAAAA TCAGCTCTAA AAAACTCAGA 16680 TARACCATTT TTTTAATTTT TTGCTTAATC ATTAATAGTG CCAATCCAAG GTTATCCACA 16740 ACAAATTTCA AATCCAATTT TGAATTTTCC TGATATACTT TTGAAATGTG TGTGTGTCCT 16800 10 GGGGATGCAA ACCAGTTTTT ATGGTAATAT ACCTAACAAA ATTTTGGAAG GCAAATCTCT 16860 TAAATACCAT GCACCTATTT CAAAACATAA TTGCAATAAT TCTGTATGCG CTTTGCTATT 16920 GGTATTTGTT TAGTTACTCC CTTCCAAGCC CTCTCTGAAT TAACAAGTTG GGTTTTATTA 16980 TGCAGATGAT ATTAACTTGA TCATCTTCTT CCTATTTCTC TGTCATGGTC AGAAGATAGG 17040 AATTGAGGTT CTTTTCCAAA TGAGGCACAG TTCTCCATGG CTATGAGACT CCATTTATGC 17100 ATCAGGAGTA AAGGGGTCTT GTGTTTTTAG GTGAGGTTCC TGGAGCAGCA GAACCAGGTA 17160 15 CTGCAACAA AATGGGAGCT GCTGCAGCAG GTAGATACCT CCACTAGAAC CCATAATTTA 17220 GAGCCCTACT TTGAGTCATT CATCAACAAT CTCCGAAGGA GAGTGGACCA ACTGAAGAGT 17280 GATCAATCTC GGTTGGATTC GGAACTGAAG AACATGCAGG ACATGGTGGA GGATTACCGG 17340 AACAAGTAAG GGACCCTGTC TGGGCAGTTC TTAACTTTTG CTGTAAAAGA GTTCCAGAAA 17400 20 GTARTARGCT ARGATCATGA AGCAGCATGT AGCTATGTCT TTTCTTAGGT TAGAGGCACA 17460 TCAGTTTGAC ATTTCAGAA ATCTTCATTT TCTCAGGAGA TGGAAATAGT CTAGTGGTTT 17520 TATTGCTCAG TAGAAAGTAG TGGCCAATAT GTCCTAGGTT CATAATAGAA AGGCAGTGAT 17580 AGGCAATGCC ACCTTTAGTT TAGAATGCTG GACTTCAGGT CTTACCACCT CTGAATCTCC 17640 TAATTGTTTC TGCTTTCCTG CAGGTATGAG GATGAAATCA ACAAGCGGAC AAATGCAGAG 17700 25 ANTGARTTG TGACCATCAA GAAGGTAAGC AAATTCTGTA GGACGGAACT CACATTTGAA 17760 ATAAATAAGG GAAGAGGGTC TCCAATTACT AAGCAGAAAG CAGCCATGAT ATGGAGAGCC 17820 AGGTAGTAGA CCTGGGGAGT ATATGGAGTG GGGCTATATT TTTCACATCA TCATGGACCT 17880 GGACTGATCC AGGCACTTGG CTTCTCCATA TTTCCCAGCA CCTTACATAG TAAGTGGAGT 17940 GGCAGATTCT CAGCAAGCCA GGCACACTCC CTTGATGGTG CTATCCGGGG GTGGGACAGT 18000 30 TGCTTGGAGA ATCCCCTCAC AGGTAATGAG AGGGACCTGC CCTGGAGAGA ACGTGCCTTC 18120 ATGATGTCCC TTGTTCCTCT AGGATGTGGA TGGTGCTTAT ATGACCAAGG TGGACCTTCA 18180 GGCCAAACTT GACAACTTGC AGCAGGAAAT TGATTTCCTT ACAGCACTCT ACCAAGCAGT 18240 AAGTCTTCCA GTTTCAACCA AGTTTATCTA AATGGAGAGT TTTTAAGCCG GAACCCACAA 18300 ² 35 CGATTCAGAA GAATAGATAT TTATCTTTTA TTTCCTGACT GCTTTCTCTG TCTAAGTTGT 18360 TTTTTGTTTT AGTGCTGTAA GAGTCACTAA CCTATTATGT CTTGCAGGAG TTGTCTCAGA 18420 tgcagactca aatcagtgaa actaatgtca tectetetat ggacaacaac egcagteteg 18480 acctggacag catcattgct gaggtcaagg cccagtacga ggatatagcc cagaagagca 18540 ARGCTGAGGC CGAGTCCTTG TACCAGAGCA AGGTGAGTGG GCTGAAACCG TAGCCAGTTT 18600

CCCTGAAATG GCTTGTCTTG CTATCCTGTG TTATCTCATG TATGTGTGCC TGTGCCATGC 18660 TGAGTTCTGC CTACATTTAA CAAACGCTAT CTACCATCTT TAGTATGAAG AGCTGCAGAT 18720 CACTGCTGGC AGACATGGGG ATAGTGTGAG AAATTCAAAG ATAGAAATTT CTGAGCTGAA 18780 TCGTGTGATC CAGAGACTTA GATCTGAAAT CGACAATGTC AAGAAGCAGG TATGTGCTTT 18840 CTCCTTCTAC CACTCAGCTG TATGGAATGG GGGTAACCCT CAGGTAAAGG GCGAGTGCTT 18900 5 TCCTAGTTTT GAATCTTGCA ATTCAGCCCA AGGCTACATT ATTAGCCCTG GTTCCTTTTC 18960 TGACTATGCT AGTTTCCAGA ATGCAGCCAT CATGCTGGGT TCTCTTTAGG GAAATCTGTG 19020 AGAATGGCCT AGTAGAGAAA GATGGGATGG TCAATGTGAG TGATCTAGCC TATGACCCAA 19080 AGTGGACTTA AGAGTTGGGG AGTGAGAGGA AGGGCAGCCA GGAGGTTTTA GAGTAGGTGT 19140 TTAGAAGAAT GTCAAGTCTG TAAGGGTTGT AGGAGCCTTG ACTCAGGGCC AAGAGAGGCT 19200 10. GTTGAGTTAT CCCTAAGGTC TTTTAAGGAA GTCAACATGG TGATGTGTTA TCTGGAGGTG 19260 GGTGTGAGAT GACTTAAGGC CAAGTGGTTC TGTTGGACTC ATTATTGGCC TCACTGGAGT 19320 GGGGAGACCA ATTGGGATGA GGAGGCCTAG TGGGGAATGC ATATTATGAG AGGGTGTCAT 19380 ATCTTTTCA GATCTCCAAC TTGCAGCAGT CCATCAGTGA TGCAGAGCAG CGTGGCGAGA 19440 ATGCCCTCAA GGATGCCAAG AACAAGCTGA ATGACCTGGA GGATGCCCTG CAGCAGGCCA 19500 15 AGGAAGACCT GGCCCGCCTG CTGCGTGACT ACCAGGAGCT GATGAACACC AAGCTGGCCC 19560 TGGATCTGGA GATTGCCACC TACAGGACCC TCCTGGAGGG AGAAGAAAGC AGGTGAGGAA 19620 GGGACGETGG GAGTCGAACC TCTTCTCATG GTCTTCCTTC CTTGCAAGCT GATTGTTGTT 19680 GAAGATGCAG CCATCTGATT GCAGCTTGTG CTGGGTATGG GGAAATGAAA AGTACACGGA 19740 20 GCAGGAGGAA GGACCTAGTT TTACTTTGGG AGCTGGAGTC CCAAGCTGTT TATTTTTTC 19800 TTCTAGGGCT GTAACATATC TAGAAAGAGC TTTGAGGTGG AGCAAATTAT TCTTTATCTG 19860 GGCTGCCTCA GATGGCAGCT GGCCTAAAGT CGGCATCTTT AGAGGGGGCC TTCATTGGCT 19920 GCAAGGCTCG TCTCGTTTAT ATGGGAATTT CTCCGTGTTT GTACTCTTGC TGAGAAAAAA 19980 TGACAGGTCT GGGAGGCCAG AGGGGATTGG ATTAAGTTTC AGATTAAGTG CATTGGAGAA 20040 GACCCAGATG GGGAAAGTCT TCAAGGTGGT GGAGCGGGGA ATGGGGAAGC GGTTTGGGAA 20100 25 GCTGGAGTGT CCTGAGGAAT TITCTTATTT TCTCCTACAG GATGTCTGGA GAATGTGCCC 20160 CGAACGTGAG TGTGTGTAAG TACAAGTCGA TTTCTCAGGG GCATGTGCAG GCTTTGTTGG 20220 GCTGGAAACG GAGTTGAGGT TGAAAATAAC TGAGCTTCCT CTTGCAGCTG TGAGCACAAG 20280 CCACACCACC ATCAGTGGAG GTGGCAGCCG AGGAGGTGGC GGCGGTGGCT ACGGCTCTGG 20340 AGGTAGCAGC TATGGCTCCG GAGGTGGTAG CTATGGTTCT GGAGGTGGCG GCGGCGGCGG 20400 30 CCGTGGCAGC TATGCCTCCG GAGGTGGCAG CTATGGCTCT GGAGGTGGCG GCGGCGGCCA 20460 TGGCAGCTAC GGCTCCGGAA GCAGCAGTGG GGGCTACAGA GGTGGCTCTG GAGGCGGCGG 20520 CGGCGCAGC TCTGGCGGCC GGGGCTCTGG CGGCGGAGC TCTGGAGGCT CCATAGGAGG 20580 CCGGGGATCC AGCTCTGGGG GTGTCAAGTC CTCTGGTGGC AGTTCCAGCG TGAAGTTTGT 20640 35 TTCTACCACT TATTCCGGAG TAACCAGATA AAGAGATGCC CTCTGTTTCA TTAGCTCTAG 20700 TTCTCCCCA GCATCACTAA CAAATATGCT TGGCAAGACC GAGGTCGATT TGTCCCAGCC 20760 TTACCGGAGA AAAGAGCTAT GGTTAGTTAC ACTAGCTCAT CCTATTCCCC CAGCTCTTTC 20820 TTTTCTGCTG TTTCCCAATG AAGTTTTCAG ATCAGTGGCA ATCTCAGTCC CCTGGCTATG 20880 ACCCTGCTTT GTTCTTTCCC TGAGAAACAG TTCAGCAGTG ACCACCC ACATGACATT 20940

TCAAAGCACC TCCTTAAGCC AGCCAGAGTA GGACCAGTTA GACCCAGGGT GTGGACAGCT 21000 CCTTAGCATC TTATCTCTGT GCTGTTTTGG TTTTGTACAT AAGGTGTAAG CAAGTTGTTT 21060 TTCTTTTGTG GAGAGGTCTT AAACTCCCCA TTTCCTTGTT TTGCTGCAAT AAACTGCATT 21120 TGAAATTCTC CATGTCTCGA TCGCCCTTGT TTACGGCACT GTCTAACCTG GATGGGTGTT 21180 TTGTGAGGTA AAAGAAGACA CTAGAGCCAC ATGGCATATG GGAAAGTCAT GCACACAAAC 21240 5 ATGAGAAAA TGCAGAGGCC AACCAGGCAA CATTTCACCA GACTGGAATC ACAGAGAGAG 21300 CAAGCACTTT CCCAGATGGT GGGGATGTCA TGGAGAAATG GAGAGACCGG GTGACAGGTT 21360 TIGITCATIT GAGAAGGCIT TCTTGAAAAG GGCAGTGAGC AAGCAGGITG GGAGGAAGAG 21420 GTGTGGCATT GAGAAGAAGG GAAAGTATTG CATGAAAAAG TAATTCTTCA CGTGGAACAG 21480 CCAGTAAGGA GGGGCATGAG TAATATAGGG TCAGCAGTTA CTGGAGCCAG AATACAGACT 21540 10 TTGGCCTGGG GAGTTCAAGA ACTAAGAGTG GTAATAGAGA GTTGGATATT CCATTTCCCT 21600 TCTCTTTTTG TGCCACCACC CAAAGCTCTG CATAATCTAA GAAGTTCCCT TGTTGACACA 21660 TAGCTCATAC TTGTGAAGTT GTACAACAGG ATAGCATAGT GGCCAGAAGC ATGGACAGTT 21720 GAACTCAGAT ATGCTTGGGT TTGAATCTTA CCATCACCAT TTACTAGTTC TGTAATACAG 21780 15 TGCAAGTTAC AGACATCTCT GCACCTCAGT TTTAGTATGT CTAAATTGGG GATGATAATG 21840 CCTTCCTTGT GGGGATAGTG TGAGGATTGA ATAAGATGAA TACACATGGC TGAGCACACA 21900 GCAAGCACTA AATAAGTGCC AGTTTTAATG ATAACGGTGA TGATGATGAT GATGATGATG 21960 ATGACGTAAC ATTGCTTGTG GGACTCCATA CAGCTCAGTA GATGCTTGCT CAAAGAAGCA 22020 AGTTACCARA ATTTTTGTAR TGGTTCTATG RACGTGARAR ARGCAGTCAR CTTCTCTGAG 22080 20 GATCAATTTC CTTAGTTTCC AATTAGGAAA AGTCTTCTTA GCTCCAGAGT CCCACAGGGC 22140 TAATGGAATA AGGAGAGGAT AGATCACACA TGTATTATGC AAACACAACT CAGGTGAGCT 22200 CTATTCTTCC TTCTCAGTTA TCCCTTCTGT AGGGACCCCA GTGTCCCCTG CTGTCTTTCT 22260 GTGTCCTGAC CGGGAAACAC AGTGTGCCTT GTCTACTCCA TCACTTGGCC AGCTGCATGC 22320 TTTCCTTTGC AGGCTTGAAG CAAAGCTGGG TCTCGGACAT TCTCAGGCAC TGACAAAGCT 22380 25 GTTTAGTTGT TGCTGGGAAA CACTGGGAAA TAGCCCTTTT GTTAAACACA CAGAAACTAG 22440 CCTTCGCCCT GAGCCAAATT CCTTAAACTC GTCTATGAAA TTCCATAACC TGACTCCTTA 22500 ACTGCAGACA TACCCAGCTA GAACATCCCT CATGTCCCTG TCCACCGTGA GAATGCTGCA 22560 CTTCACTCTG AACCTTTAGT CCTCCTTTTA AATACTGCAC ACTGATCACC CTGGTGTTTA 22620 GTGCTTTGTT TTTTGGAATC CCACCTGGCT CCATTTTGGG ATGGTTCCGG GCACTTCCCT 22680 30 ATGGAAATTC CCCTGCTGTC ACTGTCAGAG TGAGTCCAGC AGTGGGTTTA GCTGGATGAA 22740 ACACCACCAT GTCCATTTCC ATTCAGACTA ATGTCAGAAT TTGAAAGGCA CTATGGTAGA 22800 GTAGAAAGAA CAAGGAACTG TACTATTTAA AGGGCAGGCA AAGAAAAGGC ATCTATAGCT 22860 TATAAGATGT GTGGATCTTT GGATGTGACT TGGCCATCCT GAGCCTAAGT TGTCTTGTAG 22920 GAGAAATGGG AATGAGAATA TTTTCCTCTA GACATCAAGA GGAAAAGAAA TATAACGTGA 22980 35 ARACCTTTGT GARTTGTGAR TGTGTTATAC AGAGTAGCTA ARAGARTTAR ARAGGGAGTG 23040 ACAAAAAGT AAAAGGCAGC TGGCTGCTCA GGGCCTCCAT GGAGGGAAGT ACCTTGATAT 23100 GGTCACTGTG GCTCAGTGAC AGCTCTGCAG GGACAGGAAA TTGATTTGTT AGTGCACCCA 23160 AAGTTGAATC TGCTCCTGAG TACTGATTTA TGGGAACCAA ACACACAAGA GATGAAGGAT 23220 GTGTCAACCA GAATGTCCAG CATTAGCTTG TGGGGAAACA CATACTTCCA GTGACTGAAA 23280

	••••	TTATCAAGAG					23340
•	GTGGATTTAG						
	GAAGAGCTGA	CCTCTGGGGT	CAACAGATAT	AGCACCTCTT	TCCCAGGGAC	GCTACTGAAT	23460
	GAACAGTGAT	GTGTTCTTAT	ACTCTGGCCC	AGATTTTCTA	CATACTTTCT	TAGGTTACAA	23520
5	CTTTATTTAG	TCACATTICA	GTACTGGGGA	TACTCCTGTT	TATCTTCTTT	GGACTCGAGT	23580
	TTTTATGGGA	AGGTCATGAA	ACAGAGAAAA	ATACAATTTG	CAGGGAAACT	TACCAAGGCT	23640
	TGTAAGGTTA	CAAGGATTAA	ATGAAAACCC	TGTGTAAGTC	AGTATATAGT	GAAGAAGTAA	23700
	ATTGAGTTAG	ACCAAACGCC	AAAATGCATC	CGCATTAGAA	AGACGATAAA	GGAAGACTCT	23760
	GGATTCAGTT	CTGTTCAAAA	AACATTTTCT	GCACAAATAC	TATGTATGAG	GAACTGGGCG	23820
10	TTGGGGAGAT	GATGATGAGT	GAGACATGGT	TCTTGCTTTC	AGAGAGCCTA	GAGACCTGGG	23880
	TGGTAGCAAT	GGTAGAGATA	CATCCAAGAC	ACAGAAATAG	ATATACAGGA	ACACAGATGA	23940
	TTGAAAGTGA	TGCTTGGCAG	GGCTTTAAAG	AATGAATCAG	AGTTTTTCAG	GCAGACGAGG	24000
	ATCTTCAAGG	CAGAGGGAAT	CATATAGATA	AGGACATAGA	AGAGTGAAAT	TTCATGAAGT	24060
	AGTTAAGCAT	CTGAAGAAGC	ATGGAATTAG	TGACAAGAAA	TGATGCGGAA	AAGATATCCA	24120
15	GATCCAATCA	AGAAGGCCT	TGTTGGCATT	CTATGGAGTC	TGGACTTTGG	CTTCTGGGTC	24180
	ACAAGTTCTC	AGATGGGGTT	TTCATATCTA	TTATTAGACC	TACTATGTAC	TGGTCCAGTG	24240
•	GAAGGGAAAG	GGGTTGTCTT	ACTGCTAGTG	GAGTAGGAAT	TGGGTATGGA	CCACAGCTTG	24300
	TCTTGTTTCC	AAGTATTCCC	TAAGAAATCT	GGTCTGCTGA	TGGGAGATCT	ATTTATGGAA	24360
	ATGTCTTTTT	CCCTCAGGAA	TTTTATGTCG	GAAACAGCTG	TCATAGGTGA	GGAGGAACTG	24420
20	GTAAAAGTAC	TTAATAGGAG	AGTGTCATGG	TCAGATTGGT	GTTTTGGAAA	AGTCAGCCAG	24480
	GGCAGATTGG	AGAGGTCCAT	ATTGGAGGCA	GGAAGACTTA	AGAGACTATT	GCAAAGGTGA	24540
	AGACAAAAGA.	CGATAGGGAC	TTGCACTTTA	ATTCCAGCCC	TTAGAAGTAG	TAGAAGGTCA	24600
	GAAATGAGAA	TATGCATTAC	AGAGATAGTT	AGTTGCTATA	TCATTAGGAC	TTGGTGATAG	24660
	ATTGGATGAG	GATGCGGTTG	GGTGAGGCAA	AGAGGAGAGT	CCACATTCCT	GGTCTGGGTA	24720
25	GTAACAAAGA	ATCTAGCAAG	AGGGCTTGTG	GGGAAAGATG	CTGAGTTACG	TAGCAAGTGC	24780
•		TCCTTGTAAT					
		AATAAATCAT	•				
	TCCCTGGAAG	GAAAACCGGA	AGTAAATGGG	AGAAACTTGC	TCAAGTGGAC	AAAGTTTAAC	24960
•	AGACATGAAG	CATGAATTC					24979
			•				

(2) INFORMATION FOR SEQ ID NO:4:

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- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide

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(iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4: Met Ser Arg Lys Ser Tyr Lys His 5 (2) INFORMATION FOR SEQ ID NO:5: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 16 amino acids (B) TYPE: amino acid 10 (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: peptide (iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: NO (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5: 15 Met Ser Ser Val Lys Phe Val Ser Thr Thr Tyr Ser Gly Val Thr Arg

CLAIMS

What we claim is:

1. A keratin K1 vector for expression of a nucleic acid cassette in the epidermis comprising:

a 5' flanking region of the keratin K1 gene, said 5' flanking sequence including a keratin K1 promoter, a 5' transcribed but untranslated region and a first intron all in sequential and positional relationship for expression of the nucleic acid cassette;

a 3' flanking region of the keratin K1 gene containing Vitamin D_3 regulatory sequences, said 3' flanking region including a 3' transcribed but untranslated region and contiguous non-coding DNA containing the transcriptional termination region; and

a polylinker having a plurality of restriction endonuclease sites, said polylinker connecting the 5' flanking region to the 3' flanking region and said polylinker further providing a position for insertion of the nucleic acid cassette.

- 2. The keratin K1 vector of claim 1, wherein the 5' flanking region is approximately 1.2 kb and the 3' flanking region is approximately 3.9 kb.
- 3. The vector of claim 1, wherein the nucleic acid cassette includes a nucleic acid sequence coding for a protein, polypeptide or antisense RNA.
- 4. The vector of claim 1, wherein the cassette includes a nucleic acid sequence coding for an oncogene.
- 5. The vector of claim 4, wherein the oncogene is selected from the group consisting of ras, fos, myc, erb, src, sis and jun.
- 6. The vector of claim 1 wherein the cassette includes a nucleic acid sequence coding for a transforming gene.

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- 7. The vector of claim 1, wherein the restriction endonucleases are selected from the group consisting of Bam HI, Xma I, Kpn I, Not I Cla I and Bgl II.
- 8. The vector of claim 1, wherein the nucleic acid cassette contains the E6 or E7 transforming sequence of human papilloma virus.
- 9. The vector of claim 1, wherein the nucleic acid cassette contains the $TGF-\alpha$ sequence.
- 10. The vector of claim 1, further comprising an additional 5' flanking sequence from the 18 kb Eco RV fragment on the end of the vector.
- 11. A bioreactor comprising transformed epidermal cells including the vector of claim 3.
- 12. The bioreactor according to claim 11 wherein the vector includes a cassette having a nucleic acid sequence coding for a protein or polypeptide selected from the group consisting of a hormone, a growth factor, an enzyme, a drug, a tumor suppressor, a receptor, an apolipoprotein, a clotting factor, a tumor antigen, a viral antigen, an insect antigen, a bacterial antigen and a parasitic antigen.
- 13. The bioreactor of claim 12, wherein the nucleic acid sequence encodes proinsulin or insulin.
- 14. The bioreactor of claim 12, wherein the nucleic acid sequence encodes growth hormone.
- 15. The bioreactor of claim 12, wherein the nucleic acid sequence encodes insulin-like growth factor I, insulin-like growth factor II or insulin growth factor binding protein.
- 16. The bioreactor of claim 12, wherein the nucleic acid sequence encodes a clotting factor.
- 17. The bioreactor of claim 12, wherein the nucleic acid sequence encodes an epidermal growth factor (TGF-α), a dermal growth factor (PDGF) or an angiogenesis factor.

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- 18. The bioreactor of claim 12, wherein the nucleic acid sequence encodes a Type IV collagen, laminin, nidogen, or Type VII collagen.
- 19. The bioreactor of claim 12 for vaccine production, wherein the cassette includes a protein which induces an immunological response.
- 20. A method for ex vivo introduction of a keratin K1 vector into epidermal cells comprising the steps of contransfecting the vector with a selectable marker and selecting the transformed cells.
- 21. A noncoding fragment of a human keratin K1 gene containing regulatory sequences of SEQ. ID. No. 1.
- 22. A human keratin K1 vector having:
 - a 5' flanking region comprising nucleotides 1 to 1246 of SEQ. ID. No. 1;
 - a 3' flanking region comprising nucleotides 6891 to 10747 of SEQ. ID. No. 1; and
- a linker comprising nucleotide 2351 to 2376 SEQ. ID. No. 2 23. A method of making transgenic animals comprising the steps of:
- 23. A method of making transgenic animals comprising the steps of: collecting very early fertilized eggs;

inserting the vector of claim 3 into said fertilized eggs by micro-injecting the vector into pronuclei; and

transferring said injected eggs into pseudopregnant recipient females.

- 24. A transgenic animal containing the vector of claim 3 in its germ and somatic cells, wherein said vector was introduced into said animal or an ancestor of said animal at an embryonic stage and the nucleic acid cassette of said vector is only expressed in the epidermis.
- 25. The transgenic animal of claim 24, wherein the nucleic acid cassette contains a transforming gene sequence.

- 26. The transgenic animal of claim 25, wherein the oncogene is selected from the group in table 1.
- 27. The transgenic animal of claim 24 wherein the animal is a rodent.
- 28. A method of studying the origin of or treatment for cancer comprising the steps of:

making a transgenic animal by injecting an embryo with a human keratin K1 vector containing an oncogene in the nucleic acid cassette;

using the resultant animal or the progeny in studies of cancer.

- 29. The method of claim 28 wherein the nucleic acid cassette contains the fos oncogene sequence.
- 30. The method of claim 28 wherein the nucleic acid cassette contains the ras oncogene sequence.
- 31. The method of claim 28 wherein the test animal contains more than one oncogene.
- 32. A method for *in vivo* transduction of epidermal cells with a keratin K1 vector comprising the step of contacting the vector with epidermal cells for sufficient time to transform the epidermal cells.
- 33. A method for transient introduction of a nucleic acid cassette into a human, comprising the step of contacting a keratin K1 vector containing said cassette with an epidermal cell for sufficient time to transduce the cells with the vector.
- 34. A method of enhanced healing of a wound or surgical incision comprising the step of *in vivo* transduction of epidermal cells with a keratin K1 vector, wherein said vector includes a nucleic acid cassette having a nucleic acid sequence for a growth factor.
- 35. The method of claim 34, wherein the epidermal cells are transduced with a plurality of vectors and wherein the cassette of at least one vector includes the nucleic acid sequence of epidermal growth factor

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 $(TGF-\alpha)$, the cassette of at least one vector includes dermal growth factor (PDGF), the cassette of at least one vector includes the nucleic acid sequence for a matrix protein to anchor the epidermis to the dermis and the cassette of at least one vector includes the nucleic acid sequence for an angiogenesis factor.

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36. The method of claim 35, wherein the sequence for the matrix protein is selected from sequences coding for a protein selected from the group consisting of Type IV collagen, laminin, nidogen, and Type VII collagen.

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37. The method of claim 35, wherein the angiogenesis factor is selected from the group consisting of acid fibroblast growth factor, basic fibroblast growth factor and angiogenin.

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38. A method of treating skin ulcers comprising the steps of in vivo transduction of epidermal cells with a keratin K1 vector, wherein said vectors include a nucleic acid cassette having a nucleic acid sequence for a growth factor.

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39. The method of claim 38, wherein the epidermal cells are transduced with a plurality of vectors and wherein the cassette of at least one vector includes the nucleic acid sequence of epidermal growth factor (TGF-α), the cassette of at least one vector includes dermal growth factor (PDGF), the cassette of at least one vector includes the nucleic acid sequence for a matrix protein to anchor the epidermis to the dermis and the cassette of at least one vector includes the nucleic acid sequence for an angiogenesis factor.

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40. The method of claim 39, wherein the sequence for the matrix protein is selected from sequences coding for a protein selected from the group consisting of Type IV collagen, laminin, nidogen, and Type VII collagen.

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- 41. The method of claim 39, wherein the angiogenesis factor is selected from the group consisting of acid fibroblast growth factor, basic fibroblast growth factor and angiogenin.
- 42. A method of enhanced healing of a wound, surgical incision or skin ulcers in humans and animals, comprising the steps of:

ex vivo transduction of epidermal cells with a keratin K1 vector, wherein said vector includes a nucleic acid cassette having a nucleic acid sequence for a growth factor; and

transplanting said transduced epidermal cells into the animal or human to be treated.

- 43. The method of claim 42, wherein the epidermal cells are transduced with a plurality of vectors and wherein the cassette of at least one vector includes the nucleic acid sequence of epidermal growth factor (TGF-α), the cassette of at least one vector includes dermal growth factor (PDGF), the cassette of at least one vector includes the nucleic acid sequence for a matrix protein to anchor the epidermis to the dermis and the cassette of at least one vector includes the nucleic acid sequence for an angiogenesis factor.
- The method of claim 43, wherein the sequence for the matrix protein is selected from sequences coding for a protein selected from the group consisting of Type IV collagen, laminin, nidogen, and Type VII collagen.
- 45. The method of claim 43, wherein the angiogenesis factor is selected from the group consisting of acid fibroblast growth factor, basic fibroblast growth factor and angiogenin.
- 46. A method for treating psoriasis comprising the step of in vivo transduction of epidermal cells with a keratin K1 vector, wherein said vector includes a nucleic acid cassette having a nucleic acid sequence coding for a protein or polypeptide selected from the

- group consisting of TGF-\$\beta\$, a soluble form of cytokine receptor, and an antisense RNA.
- 47. The method of claim 46 wherein the cassette contains the sequence for TGF-β.
- 48. The method of claim 46 wherein the cassette contains a soluble form of cytokine receptor selected from the group consisting of IL-1, IL-6 and IL-8.
- 49. The method of claim 46, wherein the cassette contains an antisense RNA to a sequence selected from the group consisting of TGF-α, IL-1, IL-6 and IL-8.
- 50. A method for treating skin cancer comprising the step of in vivo transduction of epidermal cells with a keratin K1 vector, wherein said vector includes a nucleic acid cassette having a nucleic acid sequence coding for an antisense RNA for the E6 or E7 gene of human papilloma virus.
- 51. A method for treating skin cancer comprising the step of in vivo transduction of epidermal cells with a keratin K1 vector, wherein said vector includes a nucleic acid cassette having a nucleic acid sequence coding for the normal p53 protein.
- 52. A method for vaccination comprising the step of the *in vivo* transduction of epidermal cells with a keratin K1 vector, wherein said vector includes a nucleic acid cassette having a nucleic acid sequence coding for a protein or polypeptide which induces an immunological response.
- 53. The method of claim 52, wherein the cassette includes a sequence for a viral capsid protein.
- 54. The method of claim 53, wherein the capsid protein is from the human papilloma virus.

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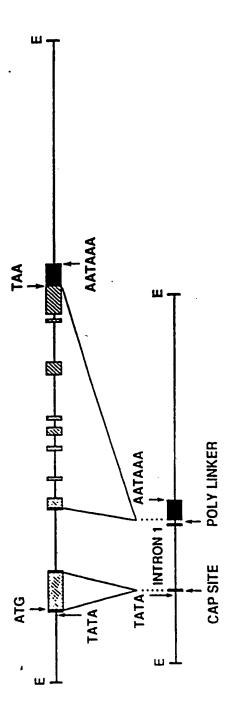


FIGURE 1

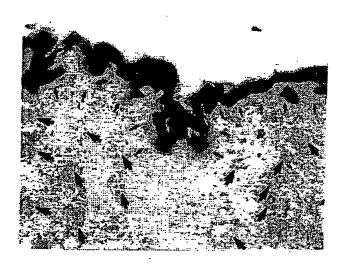


FIGURE 2A

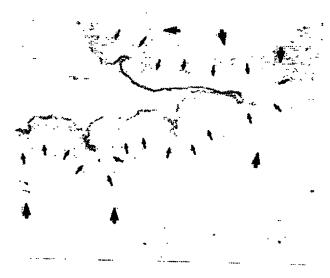


FIGURE 2B

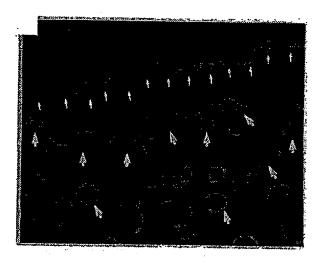
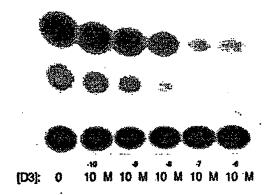


FIGURE 2C



FIGURE 2D

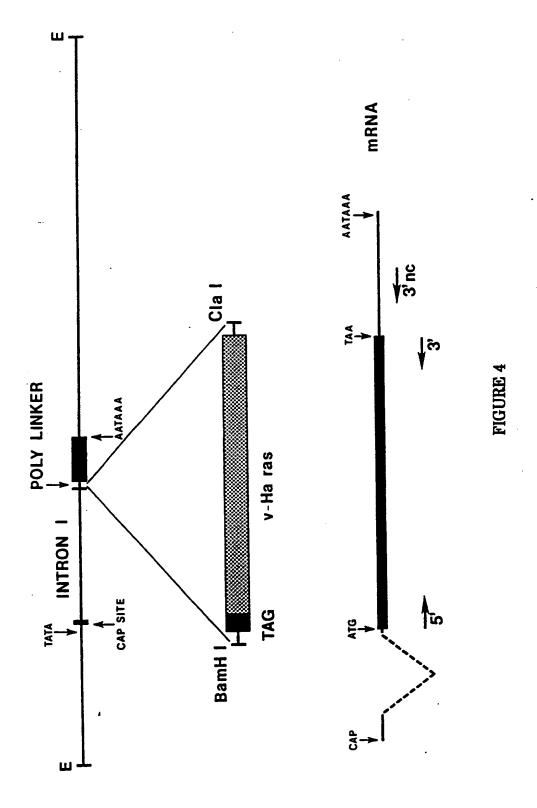
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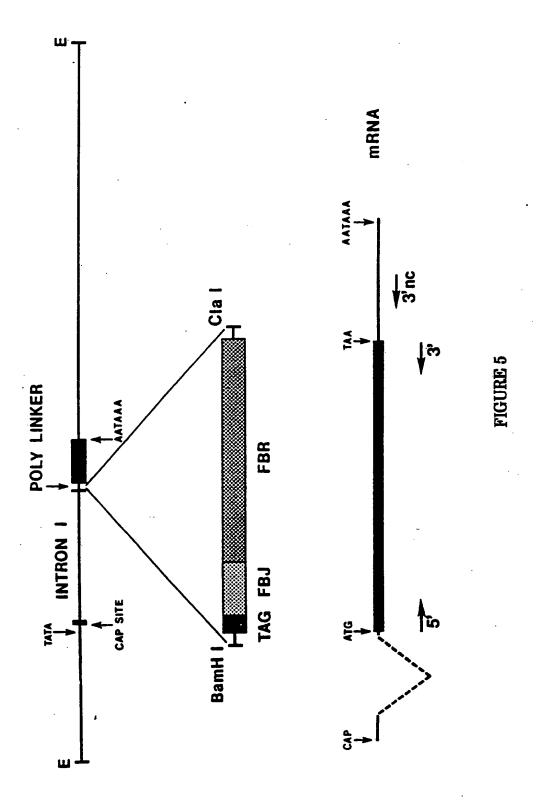
Sequence of HK1.NRE

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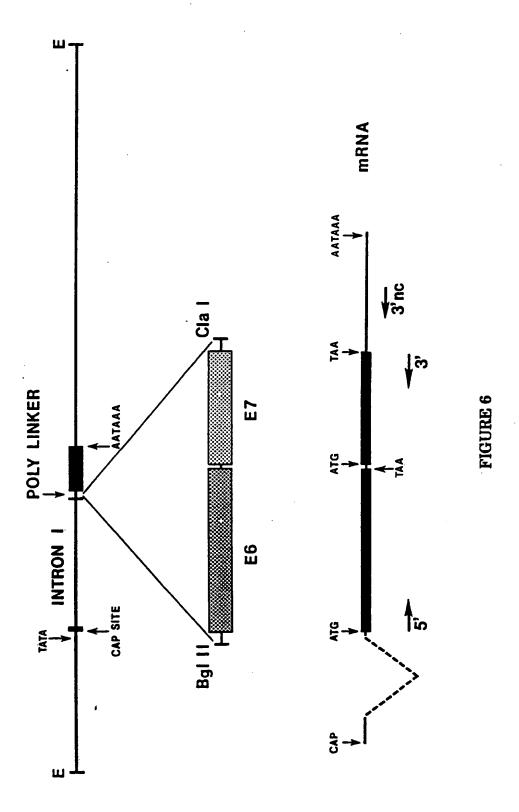
FIGURE 3



SUBSTITUTE SHEET



SUBSTITUTE SHEET



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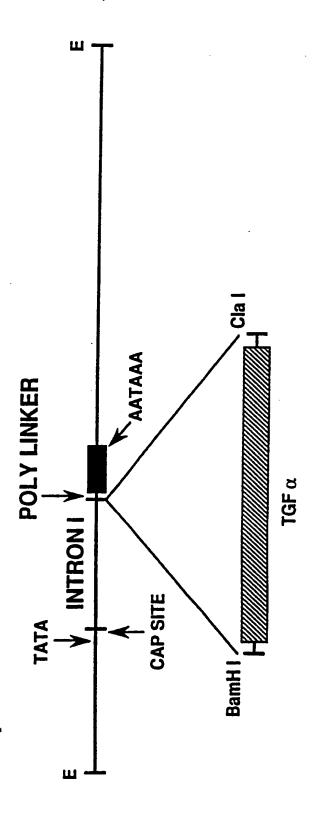


FIGURE 7

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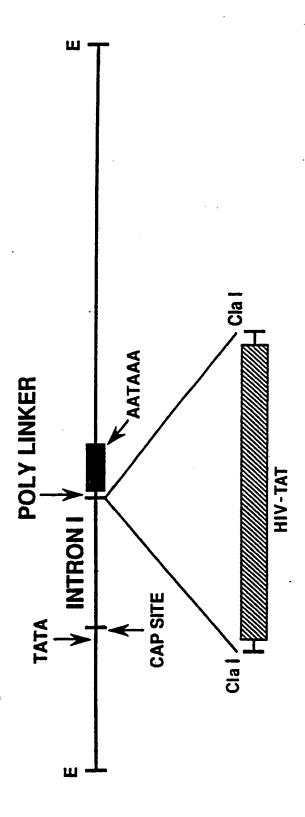
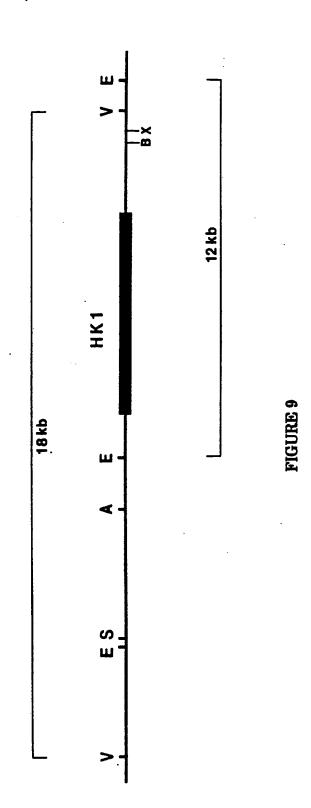


FIGURE 8

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HK1 REGULATORY ELEMENTS

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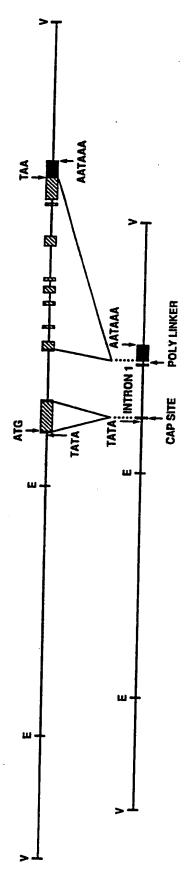


FIGURE 10

INTERNATIONAL SEARCH REPORT

International application No. PCT/US93/03985

		·	
A. CLA	ASSIFICATION OF SUBJECT MATTER: :C12N 15/00; C12N 5/00; A61K 49/00, A61K 35	/00 A61K 31/00	
US CL	:435/172.3, 240.2, 320.1, 800/2; 424/9, 93B; 514	/44	
	to International Patent Classification (IPC) or to bo	th national classification and IPC	
	documentation searched (classification system follow	and by classification symbols	
ł	435/172.3, 240.2, 320.1, 800/2; 424/9, 93B; 514/	-	•
Documenta	tion searched other than minimum documentation to	the extent that such documents are included	in the fields searched
	data base consulted during the international search (e Extra Sheet.	name of data base and, where practicable	, search terms used)
C. DOC	CUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where	appropriate, of the relevant passages	Relevant to claim No.
Y	Cell Growth & Differentiation, Volu Rosenthal et al., "A human epidermal gene is regulated by calcium but differentiation in transgenic mouse k see the entire document.	differentiation-specific keratin not negative modulators of	1-54
Y	Proc. Natl. Acad. Sci. USA, Vol. Johnson et al., "Structure of a gene fo keratin", pages 1896-1900, see the en	r the human epidermal 67-kDa	1-54
X Furthe	er documents are listed in the continuation of Box (C. See patent family annex.	
'A' doca	cial categories of cited documents: ument defining the general state of the art which is not considered e part of particular relevance	"T" later document published after the inter date and not in conflict with the applicat principle or theory underlying the inven	tion but cited to understand the
	ier document published on or after the international filing date	"X" document of particular relevance; the considered novel or cannot be consider	claimed invention cannot be ed to involve an inventive step
cited	ment which may throw doubts on priority claim(s) or which is to establish the publication date of another citation or other ial reason (as specified)	"Y" document of particular relevance; the	claimed invention cannot be
O° doct mea	ment referring to an oral disclosure, use, exhibition or other as	considered to involve an inventive a combined with one or more other such being obvious to a person skilled in the	documents, such combination
the p	ment published prior to the international filing date but later than priority date claimed	*& document member of the same patent for	
Jate of the a	ctual completion f th international search	Date of mailing f th international sear	ch report
04 JUNE 1		13 JUL 19	93
Commission Box PCT	ailing address of th ISA/US er of Patents and Trademarks D.C. 20231	JASEMINE C. CHAMBERS	yza for
	NOT APPLICABLE	Teleph ne No. (703) 308-0196	
orm PCT/IS	A/210 (second sheet)(July 1992)#		

INTERNATIONAL SEARCH REPORT

Internati nal application N . PCT/US93/03985

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	Science, Volume 217, issued 03 September 1982, Dhar et al., "Nucleotide sequence of the p21 transforming protein of Harvey murine sarcoma virus", pages 934-937, see the entire document.	3-6, 8, 9, 30
Y :	Virology, Volume 145, Issued 1985, Seedorf et al., "Human papillomavirus type 16 DNA sequence," pages 181-185, see the entire document.	3-6, 8, 9, 50, 52- 54
?	Journal of Virology, Volume 61, No. 4, Issued April 1987, Pirisi et al., "Transformation of human fibroblasts and keratinocytes with human papillomavirus type 16 DNA", pages 1061-1066, see the entire document.	11-20
	US,A, 4,736,866 (Leder et al.), 12 April 1988, see the entire document.	23-31
č	US,A, 4,497,796 (Salser et al.), 05 February 1985, see the entire document.	42-45
7	Nature, Volume 352, Issued 29 August 1991, Acsadi et al.,	32-41, 46-54
-	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document.	32-41, 40-34
- -	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire	42-45
- -	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
- -	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
- -	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
- -	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
- -	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
- -	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
- -	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
- -	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
-	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
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~	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	
-	"Human dystrophin expression in mdx mice after intramuscular injection of DNA constructs", pages 815-818, see the entire document. Science, Volume 254, Issued 06 December 1991, Dhawan et al., "Systemic delivery of human growth hormone by injection of genetically engineered myoblasts", pages 1509-1512, see the entire	

INTERNATIONAL SEARCH REPORT

Internati nal application N . PCT/US93/03985

B. FIELDS SEARCHED

Electronic data bases consulted (Name of data base and where practicable terms used):

DIALOG (Files 154, 55, 311, 312), U.S. Automated Patent System (File USPAT, 1975-1993). Search terms: keratin, K1, lone, gene, vector, transgenic, epidermis, therapy, transduction, oncogene, transforming, HPV, ras, TGF, inventors' names.

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING This ISA found multiple inventions as follows:

- I. Claims 1-22 and 42-45, drawn to a keratin K1 vector, transformed epidermal cells containing the same and a method of wound healing which comprises the ex vivo transduction of epidermal cells with a keratin K1 vector, classified in Classes 435 and 424, subclasses 320.1 and 93B, respectively, for example.
- II. Claims 23-31, drawn to a transgenic animal containing a keratin K1 vector and a method of using the same, classified in Classes 800 and 424, subclasses 2 and 9, respectively, for example.
- III. Claims 32-41 and 46-54, drawn to methods of wound healing, treating skin ulcers, treating psoriasis, treating skin cancer or vaccination, all of which comprise the <u>in vivo</u> transduction of epidermal cells with a keratin K1 vector, classified in Class 514, subclass 44, for example.

F rm PCT/ISA/210 (extra sheet)(July 1992)*